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<p>(21) International Application Number: PCT/US94/11225 (22) International Filing Date: 22 September 1994 (22.09.94) (30) Priority Data: 08/125,651 22 September 1993 (22.09.93) US 08/273,401 11 July 1994 (11.07.94) US (71) Applicant: XOMA CORPORATION [US/US]; 2910 Seventh Street, Berkeley, CA 94710 (US). (72) Inventors: COHEN, Jonathan; Dept. of Infectious Diseases & Bacteriology, Hammersmith Hospital, Du Cane Road, London W12 0NN (GB). KUNG, Ada, H., C.; Star Route, Box 48, Woodside, CA 94062 (US). LAMBERT, Lewis, H., Jr.; 45928 Omega Drive, Fremont, CA 94539 (US). LITTLE, Roger, G., II; 620 Rose Drive, Benicia, CA 94510 (US). (74) Agent: SHARP, Jeffrey, S.; Marshall, O'Toole, Gerstein, Murray & Borun, 6300 Sears Tower, 233 South Wacker Drive, Chicago, IL 60606-6402 (US).</p>		<p>(81) Designated States: AT, AU, BB, BG, BR, BY, CA, CH, CN, CZ, DE, DK, ES, FI, GB, HU, JP, KP, KR, KZ, LK, LU, LV, MG, MN, MW, NL, NO, NZ, PL, PT, RO, RU, SD, SE, SK, UA, UZ, VN, European patent (AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG).</p> <p>Published <i>With international search report. Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i></p>
<p>(54) Title: METHOD OF TREATING GRAM-NEGATIVE BACTERIAL INFECTION BY ADMINISTRATION OF BACTERICIDAL/PERMEABILITY-INCREASING (BPI) PROTEIN PRODUCT AND ANTIBIOTIC</p> <p>(57) Abstract</p> <p>The present invention relates to methods and compositions for treating gram-negative bacterial infections, using BPI protein products. Co-treatment, or concurrent administration, of BPI protein product with an antibiotic in treatment of gram-negative bacterial infections improves the therapeutic effectiveness of the antibiotic, including increasing antibiotic susceptibility of gram-negative bacteria and reversing resistance of the bacteria to antibiotics.</p>		

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**METHOD OF TREATING GRAM-NEGATIVE BACTERIAL
INFECTION BY ADMINISTRATION OF
BACTERICIDAL/PERMEABILITY-INCREASING (BPI)
PROTEIN PRODUCT AND ANTIBIOTIC**

5 This is a continuation-in-part of U.S. Application Ser. No. 08/273,401
filed July 11, 1994, which is a continuation-in-part of U.S. Application Ser. No.
08/125,651 filed September 22, 1993.

BACKGROUND OF THE INVENTION

10 The present invention relates generally to methods and compositions
for treating gram-negative bacterial infections, and more specifically to the use of
bactericidal/permeability-increasing protein (BPI) protein products for co-treatment
of such infections with an antibiotic substance. Co-treatment with BPI protein
products can improve the therapeutic effectiveness of antibiotics in gram-negative
15 bacterial infections, increase the susceptibility of gram-negative organisms to
antibiotics, and reverse resistance of gram-negative organisms to antibiotics.

20 BPI is a protein isolated from the granules of mammalian
polymorphonuclear leukocytes (PMNs or neutrophils), which are blood cells essential
in the defense against invading microorganisms. Human BPI protein has been
isolated from PMNs by acid extraction combined with either ion exchange
chromatography [Elsbach, *J. Biol. Chem.*, 254:11000 (1979)] or *E. coli* affinity
25 chromatography [Weiss, et al., *Blood*, 69:652 (1987)]. BPI obtained in such a
manner is referred to herein as natural BPI and has been shown to have potent
bactericidal activity against a broad spectrum of gram-negative bacteria. The
molecular weight of human BPI is approximately 55,000 daltons (55 kD). The amino
acid sequence of the entire human BPI protein and the nucleic acid sequence of DNA
encoding the protein have been reported in Figure 1 of Gray et al., *J. Biol. Chem.*,
264:9505 (1989), incorporated herein by reference. The Gray et al. amino acid
sequence is set out in SEQ ID NO: 69 hereto.

30 BPI is a strongly cationic protein. The N-terminal half of BPI accounts
for the high net positive charge: the C-terminal half of the molecule has a net charge
of -3. [Elsbach and Weiss (1981), *supra*.] A proteolytic N-terminal fragment of BPI

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having a molecular weight of about 25 kD has an amphipathic character, containing alternating hydrophobic and hydrophilic regions. This N-terminal fragment of human BPI possesses the anti-bacterial efficacy of the naturally-derived 55 kD human BPI holoprotein. [Ooi et al., *J. Bio. Chem.*, 262: 14891-14894 (1987)]. In contrast to the N-terminal portion, the C-terminal region of the isolated human BPI protein displays only slightly detectable anti-bacterial activity against gram-negative organisms. [Ooi et al., *J. Exp. Med.*, 174:649 (1991).] An N-terminal BPI fragment of approximately 23 kD, referred to as "rBPI₂₃," has been produced by recombinant means and also retains anti-bacterial activity against gram-negative organisms. Gazzano-Santoro et al., *Infect. Immun.* 60:4754-4761 (1992).

The bactericidal effect of BPI has been reported to be highly specific to gram-negative species, e.g., in Elsbach and Weiss, *Inflammation: Basic Principles and Clinical Correlates*, eds. Gallin et al., Chapter 30, Raven Press, Ltd. (1992). BPI is commonly thought to be non-toxic for other microorganisms, including yeast, and for higher eukaryotic cells. Elsbach and Weiss (1992), *supra*, reported that BPI exhibits anti-bacterial activity towards a broad range of gram-negative bacteria at concentrations as low as 10^{-8} to 10^{-9} M, but that 100- to 1,000-fold higher concentrations of BPI were non-toxic to all of the gram-positive bacterial species, yeasts, and higher eukaryotic cells tested at that time. It was also reported that BPI at a concentration of 10^{-4} M or 160 μ g/ml had no toxic effect, when tested at a pH of either 7.0 or 5.5, on the gram-positive organisms *Staphylococcus aureus* (four strains), *Staphylococcus epidermidis*, *Streptococcus faecalis*, *Bacillus subtilis*, *Micrococcus lysodeiktricus*, and *Listeria monocytogenes*. BPI at 10^{-4} M reportedly had no toxic effect on the fungi *Candida albicans* and *Candida parapsilosis* at pH 7.0 or 5.5, and was non-toxic to higher eukaryotic cells such as human, rabbit and sheep red blood cells and several human tumor cell lines. See also Elsbach and Weiss, *Advances in Inflammation Research*, ed. G. Weissmann, Vol. 2, pages 95-113 Raven Press (1981). This reported target cell specificity was believed to be the result of the strong attraction of BPI for lipopolysaccharide (LPS), which is unique to the outer membrane (or envelope) of gram-negative organisms.

The precise mechanism by which BPI kills gram-negative bacteria is not yet completely elucidated, but it is believed that BPI must first bind to the surface

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of the bacteria through electrostatic and hydrophobic interactions between the cationic BPI protein and negatively charged sites on LPS. LPS has been referred to as "endotoxin" because of the potent inflammatory response that it stimulates, i.e., the release of mediators by host inflammatory cells which may ultimately result in irreversible endotoxic shock. BPI binds to lipid A, reported to be the most toxic and most biologically active component of LPS.

In susceptible gram-negative bacteria, BPI binding is thought to disrupt LPS structure, leading to activation of bacterial enzymes that degrade phospholipids and peptidoglycans, altering the permeability of the cell's outer membrane, and initiating events that ultimately lead to cell death. [Elsbach and Weiss (1992), *supra*]. BPI is thought to act in two stages. The first is a sublethal stage that is characterized by immediate growth arrest, permeabilization of the outer membrane and selective activation of bacterial enzymes that hydrolyze phospholipids and peptidoglycans. Bacteria at this stage can be rescued by growth in serum albumin supplemented media [Mannion et al., *J. Clin. Invest.*, 85:853-860 (1990)]. The second stage, defined by growth inhibition that cannot be reversed by serum albumin, occurs after prolonged exposure of the bacteria to BPI and is characterized by extensive physiologic and structural changes, including apparent damage to the inner cytoplasmic membrane.

Initial binding of BPI to LPS leads to organizational changes that probably result from binding to the anionic groups in the KDO region of LPS, which normally stabilize the outer membrane through binding of Mg^{++} and Ca^{++} . Attachment of BPI to the outer membrane of gram-negative bacteria produces rapid permeabilization of the outer membrane to hydrophobic agents such as actinomycin D. Binding of BPI and subsequent gram-negative bacterial killing depends, at least in part, upon the LPS polysaccharide chain length, with long O-chain bearing, "smooth" organisms being more resistant to BPI bactericidal effects than short O-chain bearing, "rough" organisms [Weiss et al., *J. Clin. Invest.* 65: 619-628 (1980)]. This first stage of BPI action, permeabilization of the gram-negative outer envelope, is reversible upon dissociation of the BPI, a process requiring the presence of divalent cations and synthesis of new LPS [Weiss et al., *J. Immunol.* 132: 3109-3115 (1984)]. Loss of gram-negative bacterial viability, however, is not reversed by processes which restore the envelope integrity, suggesting that the bactericidal action is

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mediated by additional lesions induced in the target organism and which may be situated at the cytoplasmic membrane (Mannion et al., *J. Clin. Invest.* 86: 631-641 (1990)). Specific investigation of this possibility has shown that on a molar basis BPI is at least as inhibitory of cytoplasmic membrane vesicle function as polymyxin B (In't Veld et al., *Infection and Immunity* 56: 1203-1208 (1988)) but the exact mechanism as well as the relevance of such vesicles to studies of intact organisms has not yet been elucidated.

BPI is also capable of neutralizing the endotoxic properties of LPS to which it binds. Because of its bactericidal properties for gram-negative organisms and its ability to neutralize LPS, BPI can be utilized for the treatment of mammals suffering from diseases caused by gram-negative bacteria, such as bacteremia or sepsis.

U.S. Patent No. 5,198,541 discloses recombinant genes encoding and methods for expression of BPI proteins, including BPI-holoprotein and fragments of BPI. It also describes the use of N-terminal fragments of BPI protein for co-treatment with certain antibiotics, specifically penicillin, cephalosporins, rifampicin and actinomycin D.

Gram-negative bacteria include bacteria from the following species: *Acidaminococcus*, *Acinetobacter*, *Aeromonas*, *Alcaligenes*, *Bacteroides*, *Bordetella*, *Branhamella*, *Brucella*, *Cahymmatobacterium*, *Campylobacter*, *Cardiobacterium*, *Chromobacterium*, *Citrobacter*, *Edwardsiella*, *Enterobacter*, *Escherichia*, *Flavobacterium*, *Francisella*, *Fusobacterium*, *Haemophilus*, *Klebsiella*, *Legionella*, *Moraxella*, *Morganella*, *Neisseria*, *Pasturella*, *Plesiomonas*, *Proteus*, *Providencia*, *Pseudomonas*, *Salmonella*, *Serratia*, *Shigella*, *Streptobacillus*, *Veillonella*, *Vibrio*, and *Yersinia* species.

Antibiotics are natural chemical substances of relatively low molecular weight produced by various species of microorganisms, such as bacteria (including *Bacillus* species), actinomycetes (including *Streptomyces*) and fungi, that inhibit growth of or destroy other microorganisms. Substances of similar structure and mode of action may be synthesized chemically, or natural compounds may be modified to produce semi-synthetic antibiotics. These biosynthetic and semi-synthetic derivatives are also effective as antibiotics. The major classes of antibiotics are (1) the β -

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lactams, including the penicillins, cephalosporins and monobactams; (2) the aminoglycosides, e.g., gentamicin, tobramycin, netilmycin, and amikacin; (3) the tetracyclines; (4) the sulfonamides and trimethoprim; (5) the fluoroquinolones, e.g., ciprofloxacin, norfloxacin, and ofloxacin; (6) vancomycin; (7) the macrolides, which include for example, erythromycin, azithromycin, and clarithromycin; and (8) other antibiotics, e.g., the polymyxins, chloramphenicol and the lincosamides.

Antibiotics accomplish their anti-bacterial effect through several mechanisms of action which can be generally grouped as follows: (1) agents acting on the bacterial cell wall such as bacitracin, the cephalosporins, cycloserine, fosfomycin, the penicillins, ristocetin, and vancomycin; (2) agents affecting the cell membrane or exerting a detergent effect, such as colistin, novobiocin and polymyxins; (3) agents affecting cellular mechanisms of replication, information transfer, and protein synthesis by their effects on ribosomes, e.g., the aminoglycosides, the tetracyclines, chloramphenicol, clindamycin, cycloheximide, fucidin, lincomycin, puromycin, rifampicin, other streptomycins, and the macrolide antibiotics such as erythromycin and oleandomycin; (4) agents affecting nucleic acid metabolism, e.g., the fluoroquinolones, actinomycin, ethambutol, 5-fluorocytosine, griseofulvin, rifamycins; and (5) drugs affecting intermediary metabolism, such as the sulfonamides, trimethoprim, and the tuberculostatic agents isoniazid and para-aminosalicylic acid. Some agents may have more than one primary mechanism of action, especially at high concentrations. In addition, secondary changes in the structure or metabolism of the bacterial cell often occur after the primary effect of the antimicrobial drug.

The penicillins have a characteristic double-ring system composed of a β -lactam ring, which provides the antibacterial activity, and a thiazolidene ring. The penicillins are differentiated by a single side chain that is unique for each penicillin. The compounds are bactericidal and act by inhibiting bacterial transpeptidase, an enzyme involved in synthesis of the bacterial cell wall. Because of their mechanism of action, penicillins are generally active against growing, but not resting, cells. Penicillins, especially penicillin G, have largely gram-positive activity: the relative insensitivity of gram-negative rods to penicillin G and several other penicillins is probably due to the permeability barrier of the outer membrane of gram-

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negative bacteria. Ampicillin, carbenicillin, ticarcillin, and some other penicillins are active against gram-negative bacteria because they can pass through this outer membrane. Penicillins have relatively few adverse effects, the most important of which are the hypersensitivity (allergic) reactions. These compounds are widely distributed in the body, but do not enter cells and do not usually accumulate in CSF.

Bacterial resistance to the penicillins is by production of the enzyme β -lactamase, which catalyzes hydrolysis of the β -lactam ring. The percentage of bacteria resistant to penicillin has risen to about 80%. Several penicillins, including methicillin, oxacillin, cloxacillin, dicloxacillin and nafcillin, are not affected by the β -lactamase of staphylococci. These antibiotics are useful against most β -lactamase-producing species of *Staphylococcus*. However, a small number of species are resistant even to these penicillins. Some penicillins, amoxicillin and ticarcillin, are marketed in combination with clavulanic acid, which is a β -lactamase inhibitor that covalently binds to the enzyme and prevents it from hydrolyzing the antibiotics. Another inhibitor, sulbactam, is marketed in combination with ampicillin.

The cephalosporins are characterized by a β -lactam ring, like the penicillins, but have an adjacent dihydrothiazine ring instead of a thiazolidene ring. For convenience, these compounds are generally classified by generations. The first generation includes cephalothin, cephapirin, cefazolin, cephalexin, cephradine and cefadroxil. These drugs generally have excellent gram-positive activity except for enterococci and methicillin-resistant staphylococci, and have only modest gram-negative coverage. The second generation includes cefamandole, cefoxitin, ceforanide, cefuroxime, cefuroxime axetil, cefaclor, cefonicid and cefotetan. This generation generally loses some gram-positive activity by weight and gains limited gram-negative coverage. The third generation includes cefotaxime, moxalactam, ceftizoxime, ceftriaxone, cefoperazone and ceftazidime. These compounds generally sacrifice further gram-positive activity by weight but gain substantial gram-negative coverage against *Enterobacter* and sometimes are active against *Pseudomonas*. The cephalosporins bind to penicillin-binding proteins with varying affinity. Once binding occurs, protein synthesis is inhibited. Cephalosporins are usually well tolerated; adverse effects include hypersensitivity reactions and gastrointestinal effects. Cephalosporins may interact with nephrotoxic drugs, particularly aminoglycosides.

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to increase toxicity. Resistance to cephalosporins is mediated by several mechanisms, including production of β -lactamase, although some strains that do not produce β -lactamase are nevertheless resistant.

5 Imipenem is a N-formimidoyl derivative of the mold product thienamycin. It contains a β -lactam ring and somewhat resembles penicillin except for differences in the second ring. It has activity against both gram-positive and gram-negative organisms and is resistant to most β -lactamases, although not those from *Pseudomonas*. It is marketed in combination with cilastin, a compound that inhibits
10 inactivation of imipenem in the kidney by renal dihydropeptidase I enzyme. Cilastin increases the concentration of imipenem in urine, although not in blood.

Aztreonam is the first of a new group of antibiotics referred to as the monobactams. These agents have a β -lactam ring but lack the second ring characteristic of the penicillins and cephalosporins. It acts by binding to penicillin-binding proteins, and produces long, filamentous bacterial shapes that eventually lyse.
15 Aztreonam is active only against aerobic gram-negative bacteria, is susceptible to inactivation by some β -lactamases, and has few adverse effects.

The aminoglycosides contain amino sugars linked to an aminocyclitol ring by glycosidic bonds. They have similar mechanisms of action and properties, but differ somewhat in spectrum of action, toxicity, and susceptibility to bacterial resistance. The compounds are bactericidal, with activity against both gram-positive
20 and gram-negative organisms, and act by binding to proteins on the 30S ribosome of bacteria and inhibiting protein synthesis. The aminoglycosides also bind to isolated LPS and have a very weak outer membrane permeabilizing effect. [Taber et al., *Microbiological Reviews* 53: 439-457 (1987); Kadurugamuwa et al., *Antimicrobial Agent and Chemotherapy*, 37: 715-721 (1993); Vaara, *Microbiological Reviews* 56: 395-411 (1992)]. This class of antibiotics includes amikacin, gentamicin, kanamycin,
25 neomycin, netilmycin, paromomycin and tobramycin. The aminoglycosides are usually reserved for more serious infections because of severe adverse effects including ototoxicity and nephrotoxicity. There is a narrow therapeutic window between the concentration required to produce a therapeutic effect, e.g., 8 μ g/ml for
30 gentamicin, and the concentration that produces a toxic effect, e.g., 12 μ g/ml for

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gentamicin. Neomycin in particular is highly toxic and is never administered parenterally.

5 Tetracyclines have a common four-ring structure and are closely congeneric derivatives of the polycyclic naphthacenecarboxamide. The compounds are bacteriostatic, and inhibit protein synthesis by binding to the 30S subunit of microbial ribosomes and interfering with attachment of aminoacyl tRNA. The compounds have some activity against both gram-positive and gram-negative bacteria; however, their use is limited because many species are now relatively resistant. Adverse effects include gastrointestinal effects, hepatotoxicity with large doses, and
10 nephrotoxicity in some patients. This antibiotic class includes tetracycline, chlortetracycline, demeclocycline, doxycycline, methacycline, minocycline and oxytetracycline.

The sulfonamides are derivatives of sulfanilamide, a compound similar in structure to para-aminobenzoic acid (PABA), which is an essential precursor for
15 bacterial synthesis of folic acid. The compounds are generally bacteriostatic, and act by competitively inhibiting incorporation of PABA into tetrahydrofolic acid, which is a required cofactor in the synthesis of thymidines, purines and DNA. Sulfonamides have a wide range of activity against gram-positive and gram-negative bacteria, but their usefulness has diminished with increasingly high prevalence of bacterial
20 resistance. The sulfonamide class of antibiotics includes sulfacytine, sulfadiazine, sulfamethizole, sulfisoxazole, sulfamethoxazole, sulfabenzamide and sulfacetamide. Adverse effects include hypersensitivity reactions and occasional hematological toxicity.

Trimethoprim is an inhibitor of the dihydrofolate reductase enzyme,
25 which converts dihydrofolic to tetrahydrofolic acid, a required factor for DNA synthesis. Adverse effects include gastrointestinal distress and rare hematological toxicity. Trimethoprim is also available in combination with sulfamethoxazole (also known as co-trimoxazole). The combination is usually bactericidal, although each agent singly is usually bacteriostatic. The combination is the drug of choice for
30 *Salmonella* infections, some *Shigella* infections, *E. coli* traveler's diarrhea and *Pneumocystis carinii* pneumonia.

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The fluoroquinolones and quinolones are derivatives of nalidixic acid, a naphthyridine derivative. These compounds are bactericidal, and impair DNA replication, transcription and repair by binding to the DNA and interfering with DNA gyrase, an enzyme which catalyzes negative supercoiling of DNA. The fluoroquinolones, which include norfloxacin, ciprofloxacin, and ofloxacin, and the quinolones, which include cinoxacin, have a broad spectrum of antimicrobial activity against gram-negative and gram-positive organisms. These compounds distribute widely through extravascular tissue sites, have a long serum half-life, and present few adverse effects. Because of their effect on DNA, the drugs are contraindicated in pregnant patients and in children whose skeletal growth is incomplete.

Vancomycin is a glycopeptide, with a molecular weight of about 1500, produced by a fungus. It is primarily active against gram-positive bacteria. The drug inhibits one of the final steps in synthesis of the bacterial cell wall, and is thus effective only against growing organisms. It is used to treat serious infections due to gram-positive cocci when penicillin G is not useful because of bacterial resistance or patient allergies. Vancomycin has two major adverse effects, ototoxicity and nephrotoxicity. These toxicities can be potentiated by concurrent administration of another drug with the same adverse effect, such as an aminoglycoside.

The macrolides are bacteriostatic and act by binding to the 50S subunit of 70S ribosomes, resulting in inhibition of protein synthesis. They have a broad spectrum of activity against gram-positive and gram-negative bacteria and may be bacteriostatic or bactericidal, depending on the concentration achieved at sites of infection. The compounds distribute widely in body fluids. Adverse effects include gastrointestinal distress and rare hypersensitivity reactions. The most common macrolide used is erythromycin, but the class includes other compounds such as clarithromycin and azithromycin.

The polymyxins are a group of closely related antibiotic substances produced by strains of *Bacillus polymyxa*. These drugs, which are cationic detergents, are relatively simple, basic peptides with molecular weights of about 1000. Their antimicrobial activity is restricted to gram-negative bacteria. They interact strongly with phospholipids and act by penetrating into and disrupting the structure of cell membranes. Polymyxin B also binds to the lipid A portion of

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endotoxin and neutralizes the toxic effects of this molecule. Polymyxin B has severe adverse effects, including nephrotoxicity and neurotoxicity, and should not be administered concurrently with other nephrotoxic or neurotoxic drugs. The drug thus has limited use as a therapeutic agent because of high systemic toxicity, but may be used for severe infections, such as *Pseudomonas aeruginosa* meningitis, that respond poorly to other antibiotics.

Chloramphenicol inhibits protein synthesis by binding to the 50S ribosomal subunit and preventing binding of aminoacyl tRNA. It has a fairly wide spectrum of antimicrobial activity, but is only reserved for serious infections, such as meningitis, typhus, typhoid fever, and Rocky Mountain spotted fever, because of its severe and fatal adverse hematological effects. It is primarily bacteriostatic, although it may be bactericidal to certain species.

Lincomycin and clindamycin are lincosamide antimicrobials. They consist of an amino acid linked to an amino sugar. Both inhibit protein synthesis by binding to the 50S ribosomal subunit. They compete with erythromycin and chloramphenicol for the same binding site but in an overlapping fashion. They may be bacteriostatic or bactericidal, depending on relative concentration and susceptibility. Gastrointestinal distress is the most common side effect. Other adverse reactions include cutaneous hypersensitivity, transient hematological abnormalities, and minor elevations of hepatic enzymes. Clindamycin is often the drug of choice for infections caused by anaerobic bacteria or mixed aerobic/anaerobic infections, and can also be used for susceptible aerobic gram-positive cocci.

Some drugs, e.g. aminoglycosides, have a small therapeutic window. For example, 2 to 4 $\mu\text{g/ml}$ of gentamicin or tobramycin may be required for inhibition of bacterial growth, but peak concentrations in plasma above 6 to 10 $\mu\text{g/ml}$ may result in ototoxicity or nephrotoxicity. These agents are more difficult to administer because the ratio of toxic to therapeutic concentrations is very low. Antimicrobial agents that have toxic effects on the kidneys and that are also eliminated primarily by the kidneys, such as the aminoglycosides or vancomycin, require particular caution because reduced elimination can lead to increased plasma concentrations, which in turn may cause increased toxicity. Doses of antimicrobial agents that are eliminated by the kidneys must be reduced in patients with impaired

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renal function. Similarly, dosages of drugs that are metabolized or excreted by the liver, such as erythromycin, chloramphenicol, or clindamycin, must be reduced in patients with decreased hepatic function.

5 Antibiotic resistance in bacteria is an increasingly troublesome problem. The accelerating development of antibiotic-resistant bacteria, intensified by the widespread use of antibiotics in farm animals and overprescription of antibiotics by physicians, has been accompanied by declining research into new antibiotics with different modes of action. [*Science*, 264: 360-374 (1994).] Antibiotic resistance, once acquired, can be rapidly spread to other bacteria, including bacteria of a
10 different species. There are some species of bacteria that are resistant to all but one antibiotic; it may be only a matter of time before the appearance of bacterial strains that are resistant to all antibiotics.

Bacteria acquire resistance to antibiotics through several mechanisms:
15 (1) production of enzymes that destroy or inactivate the antibiotic [Davies, *Science*, 264:375-381 (1994)]; (2) synthesis of new or altered target sites on or within the cell that are not recognized by the antibiotic [Spratt, *Science*, 264:388-393 (1994)]; (3) low permeability to antibiotics, which can be reduced even further by altering cell wall proteins, thus restricting access of antibiotics to the bacterial cytoplasmic machinery; (4) reduced intracellular transport of the drug; and (5) increased removal
20 of antibiotics from the cell via membrane-associated pumps [Nikaido, *Science*, 264:382-387 (1994)].

The susceptibility of a bacterial species to an antibiotic is generally determined by two microbiological methods. A rapid but crude procedure uses commercially available filter paper disks that have been impregnated with a specific
25 quantity of the antibiotic drug. These disks are placed on the surface of agar plates that have been streaked with a culture of the organism being tested, and the plates are observed for zones of growth inhibition. A more accurate technique, the broth dilution susceptibility test, involves preparing test tubes containing serial dilutions of the drug in liquid culture media, then inoculating the organism being tested into the
30 tubes. The lowest concentration of drug that inhibits growth of the bacteria after a suitable period of incubation is reported as the minimum inhibitory concentration.

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The resistance or susceptibility of an organism to an antibiotic is determined on the basis of clinical outcome, i.e., whether administration of that antibiotic to a subject infected by that organism will successfully cure the subject. While an organism may literally be susceptible to a high concentration of an antibiotic *in vitro*, the organism may in fact be resistant to that antibiotic at physiologically realistic concentrations. If the concentration of drug required to inhibit growth of or kill the organism is greater than the concentration that can safely be achieved without toxicity to the subject, the microorganism is considered to be resistant to the antibiotic. To facilitate the identification of antibiotic resistance or susceptibility using *in vitro* test results, the National Committee for Clinical Laboratory Standards (NCCLS) has formulated standards for antibiotic susceptibility that correlate clinical outcome to *in vitro* determinations of the minimum inhibitory concentration of antibiotic.

Thus, there exists a desire in the art for agents that could act as adjuncts to conventional antibiotic therapy and that could act to improve the therapeutic effectiveness of antibiotics.

SUMMARY OF THE INVENTION

The present invention generally provides methods and compositions for prophylaxis or treatment of gram-negative bacterial infections, using BPI protein products. The methods and compositions, in addition to being useful for treatment, are useful for prophylaxis of patients at high risk of gram-negative bacterial infection, e.g., patients who will undergo abdominal or genitourinary surgery, or trauma victims. Specifically, the present invention provides, in a method for treating a gram-negative bacterial infection with an antibiotic, the improvement comprising the step of concurrently administering BPI protein product in an amount effective to improve the therapeutic effectiveness of the antibiotic.

The present invention is based upon the finding that BPI protein product is useful as adjunct therapy with conventional antibiotics, and specifically the finding that concurrent administration, or co-treatment, of a BPI protein product and an antibiotic or combination of antibiotics can improve the therapeutic effectiveness of the antibiotic or combination of antibiotics. BPI protein product may improve the

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therapeutic effectiveness of antibiotics in a variety of ways, including by increasing susceptibility of gram-negative bacteria to a reduced dosage of antibiotics, by effectively reversing resistance of gram-negative bacteria to antibiotics, by providing synergistic or potentiating effects beyond the individual or additive effects of the BPI protein product or antibiotic alone, or by neutralizing endotoxin released by bacteria killed by antibiotics. Concurrent administration of BPI protein product and antibiotic provides unexpectedly superior therapeutic effects *in vivo* than either agent provides when administered alone. Concurrent administration of BPI protein product according to this improved method of treatment is effective even when the gram-negative bacteria involved are considered to be resistant to the bactericidal effects of BPI protein product alone and/or antibiotic alone.

The present invention provides a use of a BPI protein product for the manufacture of a medicament for the co-treatment with an antibiotic of a gram-negative bacterial infection in mammals. This aspect of the invention contemplates co-treatment with any antibiotic or combinations of antibiotics, including β -lactam antibiotics with and without β -lactamase inhibitors, aminoglycosides, tetracyclines, sulfonamides and trimethoprim, vancomycin, macrolides, fluoroquinolones and quinolones, polymyxins and other antibiotics.

This aspect of the invention also provides the use of a BPI protein product for the manufacture of a medicament for improving the therapeutic effectiveness of antibiotics in a gram-negative bacterial infection, use of a BPI protein product for the manufacture of a medicament for increasing the susceptibility to an antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection, and use of a BPI protein product for the manufacture of a medicament for reversing resistance to an antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

The invention utilizes any of the large variety of BPI protein products known to the art including natural BPI protein, recombinant BPI protein, BPI fragments, BPI analogs, BPI variants, and BPI peptides. Concurrent administration of BPI protein product with any antibiotic or combination of antibiotics is contemplated, including β -lactam antibiotics with or without β -lactamase inhibitors.

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aminoglycosides, tetracyclines, sulfonamides and trimethoprim, vancomycin, macrolides, fluoroquinolones and quinolones, polymyxins, and other antibiotics.

5 Either the BPI protein product or the antibiotic may be administered systemically or topically to a subject suffering from a suspected or confirmed gram-negative bacterial infection. The BPI protein product and antibiotic may be administered by different routes and may be administered simultaneously or sequentially.

10 The invention also provides pharmaceutical compositions for treatment of gram-negative bacterial infections, comprising an antibiotic and a BPI protein product in an amount effective to improve the therapeutic effectiveness of the antibiotic. Such compositions optionally comprise pharmaceutically acceptable diluents, adjuvants or carriers. The compositions may be formulated for systemic or topical administration to subjects. In addition, compositions comprising BPI protein product and an antibiotic can be used in a variety of *in vitro* uses, such as use as a bactericide to decontaminate fluids and surfaces and to sterilize surgical and other medical equipment and implantable devices, including prosthetic joints.

15 Numerous additional aspects and advantages of the invention will become apparent to those skilled in the art upon consideration of the following detailed description of the invention which describes presently preferred embodiments thereof.

BRIEF DESCRIPTION OF THE DRAWINGS

20 Figure 1 depicts survival data after treatment with rBPI₂₁ and cefamandole, separately or in combination, in an *E. coli* O111:B4 mouse peritonitis assay.

25 Figures 2-4 relate to results from an *E. coli* O111:B4 mouse peritonitis assay with rBPI₂₁ and cefamandole, separately or in combination. Figure 2 depicts survival data; Figure 3 shows bacterial counts from peritoneal lavage fluid; and Figure 4 shows bacterial counts in blood.

Figures 5A and 5B display results from two trials of treatment with

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rBPI₁₁ and cefamandole, separately or in combination, in an *E. coli* O7:K1 mouse peritonitis assay.

Figures 6-16 relate to results, including cardiovascular and metabolic findings, from an *E. coli* O7:K1 rabbit bacteremia assay with rBPI₁₁ and cefamandole, separately or in combination. Figure 6 depicts survival data; Figure 7 shows bacterial count in blood, which Figure 8 displays as percentage of bacterial dose; Figure 9 shows blood endotoxin levels; Figures 10 shows mean arterial blood pressure; Figure 11, cardiac index; Figure 12, total peripheral resistance; Figure 13, arterial oxygen tension; Figure 14, alveolar-arterial oxygen gradient; Figure 15, respiration rate; and Figure 16, arterial blood pH.

Figure 17 depicts survival data after treatment with rBPI₁₁ and gentamicin, separately or in combination, in an *E. coli* O7:K1 mouse peritonitis assay.

Figure 18 shows the bactericidal effect of rBPI₁₁ and ceftriaxone, separately or in combination, on growth of ceftriaxone-resistant *E. coli*.

Figures 19-25 relate to the synergistic effect of rBPI₁₁ on antibiotic killing curves for selected gram-negative organisms. Figure 19 shows the bactericidal effect of rBPI₁₁ alone. Figure 20 shows the effect of rBPI₁₁ in combination with trimethoprim/sulfamethoxazole; Figure 21, rBPI₁₁ with ciprofloxacin; Figure 22, rBPI₁₁ with piperacillin; Figure 23, rBPI₁₁ with cefotaxime; Figure 24, rBPI₁₁ with cefuroxime; and Figure 25, rBPI₁₁ with amikacin.

DETAILED DESCRIPTION

The present invention relates to methods and compositions for treating a gram-negative bacterial infection, using a BPI protein product. The invention is based on the unexpected discovery that, when treating a gram-negative bacterial infection with an antibiotic, the concurrent administration of BPI protein product with the antibiotic improves the therapeutic effectiveness of the antibiotic, even at doses at which the BPI protein product alone or antibiotic alone may be inactive. BPI protein product by itself typically has an antibacterial potency less than that of conventional antibiotics. However, because its administration unexpectedly improves the therapeutic effectiveness of conventional antibiotic therapy, BPI protein product

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is useful as adjunct therapy with conventional antibiotic therapy for the treatment of gram-negative bacterial infections.

"Gram-negative bacterial infection," as used herein, encompasses conditions associated with or resulting from gram-negative bacterial infection (e.g., sequelae). These conditions include gram-negative sepsis, endotoxin-related hypotension and shock, and one or more of the conditions associated therewith, including fever, metabolic acidosis, disseminated intravascular coagulation and related clotting disorders, anemia, thrombocytopenia, leukopenia, adult respiratory distress syndrome and related pulmonary disorders, renal failure and related renal disorders, hepatobiliary disease and central nervous system disorders. These conditions also include translocation of bacteria from the intestines and concomitant release of endotoxin.

BPI protein product may improve the therapeutic effectiveness of the antibiotic in a variety of ways, including by increasing susceptibility of gram-negative bacteria to a reduced dosage of antibiotics, by effectively reversing resistance of gram-negative bacteria to antibiotics, by providing synergistic or potentiating effects beyond the individual or additive effects of the BPI protein product or antibiotic alone, or by neutralizing endotoxin released by bacteria killed by antibiotics. Concurrent administration of BPI protein product and antibiotic is expected to provide more effective treatment of gram-negative bacterial infection. Concurrent administration of the two agents may provide greater therapeutic effects *in vivo* than either agent provides when administered singly. It may permit a reduction in the dosage of one or both agents with achievement of a similar therapeutic effect. Alternatively, the concurrent administration may produce a more rapid or complete bactericidal/bacteriostatic effect than could be achieved with either agent alone.

Therapeutic effectiveness is based on a successful clinical outcome, and does not require that the antimicrobial agent or agents kill 100% of the organisms involved in the infection. Success depends on achieving a level of antibacterial activity at the site of infection that is sufficient to inhibit the bacteria in a manner that tips the balance in favor of the host. When host defenses are maximally effective, the antibacterial effect required may be minimal. Reducing organism load by even one log (a factor of 10) may permit the host's own defenses to control the infection.

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In addition, augmenting an early bactericidal/bacteriostatic effect can be more important than long-term bactericidal/bacteriostatic effect. These early events are a significant and critical part of therapeutic success, because they allow time for host defense mechanisms to activate. Increasing the bactericidal rate may be particularly important for infections such as meningitis, bone or joint infections. [Stratton, *Antibiotics in Laboratory Medicine*, 3rd ed. (Lorian, V., Ed.) pp. 849-879, Williams and Wilkins, Baltimore MD (1991)].

The effect of BPI protein product to improve the therapeutic effectiveness of antibiotics *in vivo* may be demonstrated in *in vivo* animal models, or may be predicted on the basis of a variety of *in vitro* tests, including (1) determinations of the minimum inhibitory concentration (MIC) of an antibiotic required to inhibit growth of a gram-negative organism for 24 hours, (2) determinations of the effect of an antibiotic on the kinetic growth curve of a gram-negative organism, and (3) checkerboard assays of the MIC of serial dilutions of antibiotic alone or in combination with serial dilutions of BPI protein product. Exemplary models or tests are described in Eliopoulos and Moellering In *Antibiotics in Laboratory Medicine*, 3rd ed. (Lorian, V., Ed.) pp. 432-492, Williams and Wilkins, Baltimore MD (1991).

Using *in vitro* determinations of antibiotic MIC at 24 hours, a BPI protein product may be shown to reduce the MIC of the antibiotic. With this result, it is expected that concurrent administration of the BPI protein product *in vivo* will increase susceptibility of the gram-negative organism to the antibiotic. A BPI protein product may also be shown to reduce the MIC of an antibiotic from the range in which the organism is considered clinically resistant to a range in which the organism is considered clinically susceptible. With this result, it is expected that concurrent administration *in vivo* of the BPI protein product with the antibiotic will reverse resistance and effectively convert the antibiotic-resistant organism into an antibiotic-susceptible organism.

By measuring the effect of antibiotics on the *in vitro* growth curves of gram-negative organisms, in the presence or absence of a BPI protein product, the BPI protein product may be shown to enhance the early antibacterial effect of

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antibiotics at 0-24 hours. Enhancement of early bactericidal/growth inhibitory effects is important in determining therapeutic outcome.

The BPI protein product and antibiotic may also be shown to have synergistic or potentiating effects beyond the individual effects of each agent alone or the additive effects of the agents together. In a checkerboard assay, the combination of BPI protein product with antibiotics may be shown to result in a "synergistic" fractional inhibitory concentration index (FIC). The checkerboard method is based on additivity, which assumes that the result observed with multiple drugs is the sum of the separate effects of the drugs being tested; according to this system a FIC of less than 0.5 is scored as synergy, 1 is scored as additive, and greater than 1 but less than 2 is scored as indifferent. In contrast, kinetic assays are based on the idea that only one metabolic pathway at a time can be growth rate-limiting for an organism; according to this system, the combined effect of drugs that do not interact with one another (autonomous or indifferent) is simply the effect of the most active drug alone.

Concurrent administration of BPI protein products and antibiotics is shown herein to lower MICs of a variety of antibiotics for a variety of gram-negative organisms. It is also shown to reverse resistance of a variety of gram-negative organisms to antibiotics. In some cases where BPI protein product does not affect the MIC of antibiotic at 24 hours, BPI protein product is shown herein to enhance the early bactericidal effect of antibiotics on growth curves at 0-7 or 7-24 hours. The BPI protein products exert these effects even on gram-negative organisms that are not considered susceptible to the direct bactericidal or growth inhibitory effects of BPI protein product alone. It is also shown herein that the concurrent administration of BPI protein products with antibiotics *in vivo* allows a reduction in the dosages of both agents to amounts that, if administered alone, would be insufficient to exert the same clinical effect.

Either the BPI protein product or the antibiotic, or both, may be administered at levels at which neither would alone be therapeutically effective against a gram-negative bacterial infection. Alternatively, according to a preferred method, the antibiotic and BPI protein product can be administered in amounts where each would alone be therapeutically effective against a gram-negative bacterial infection

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but wherein the combination of the two antibiotics provides even more potent effects. The BPI protein product may be administered in an amount which increases susceptibility of gram-negative bacteria to reduced antibiotic dosage, or in an amount which reverses resistance of the gram-negative bacteria to an antibiotic.

5 BPI protein product is thought to interact with a variety of host defense elements present in whole blood or serum, including complement, p15 and LBP, and other cells and components of the immune system. Such interactions may result in potentiation of the activities of BPI protein product. Because of these interactions, BPI protein products can be expected to exert even greater activity *in vivo* than *in*
10 *vitro*. Thus, while *in vitro* tests are predictive of *in vivo* utility, absence of activity *in vitro* does not necessarily indicate absence of activity *in vivo*. For example, BPI has been observed to display a greater bactericidal effect on gram-negative bacteria in whole blood or plasma assays than in assays using conventional media. [Weiss et al., *J. Clin. Invest.* 90:1122-1130 (1992)]. This may be because conventional *in vitro*
15 systems lack the blood elements that facilitate or potentiate BPI's function *in vivo*, or because conventional media contain higher than physiological concentrations of magnesium and calcium, which are typically inhibitors of the antibacterial activity of BPI protein products. Furthermore, in the host, BPI protein product is available to neutralize endotoxin released during antibiotic killing of bacteria, a further clinical
20 benefit not seen in or predicted by *in vitro* tests.

It is also contemplated that the BPI protein product be administered with other products that potentiate the bactericidal activity of BPI protein products. For example, serum complement potentiates the gram-negative bactericidal activity of BPI protein products; the combination of BPI protein product and serum
25 complement provides synergistic bactericidal/growth inhibitory effects. See, e.g., Ooi et al. *J. Biol. Chem.*, 265: 15956 (1990) and Levy et al. *J. Biol. Chem.*, 268: 6038-6083 (1993) which address naturally-occurring 15 kD proteins potentiating BPI antibacterial activity. See also co-owned, co-pending PCT Application No. US94/07834 filed July 13, 1994, which corresponds to U.S. Patent Application Serial
30 No. 08/274,303 filed July 11, 1994 as a continuation-in-part of U.S. Patent Application Serial No. 08/093,201 filed July 14, 1993. These

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applications, which are all incorporated herein by reference, describe methods for potentiating gram-negative bactericidal activity of BPI protein products by administering lipopolysaccharide binding protein (LBP) and LBP protein products. LBP protein derivatives and derivative hybrids which lack CD-14 immunostimulatory properties are described in PCT Application No. US94/06931 filed June 17, 1994, which corresponds to co-owned, co-pending U.S. Patent Application Serial No. 08/261,660, filed June 17, 1994 as a continuation-in-part of U.S. Patent Application Serial No. 08/079,510, filed June 17, 1993, the disclosures of all of which are hereby incorporated by reference.

An advantage provided by the present invention is the ability to provide more effective treatment of gram-negative bacterial infection by virtue of the improved therapeutic effectiveness of the antibiotic treatment. Another advantage is the ability to treat gram-negative organisms that are normally resistant to one or more antibiotics. Yet another advantage is the ability to accelerate the killing of gram-negative organisms by antibiotics. An additional advantage is the ability to neutralize endotoxin released during antibiotic killing of bacteria. A further advantage is the ability to use lower concentrations of toxic antibiotics such as gentamicin and polymyxin B, or expensive antibiotics such as vancomycin. Because the use of some antibiotics is limited by their systemic toxicity or prohibitive cost, lowering the concentration of antibiotic required for therapeutic effectiveness reduces toxicity and/or cost of treatment, and thus allows wider use of the antibiotic. The present invention may also provide quality of life benefits due to, e.g., decreased duration of therapy, reduced stay in intensive care units or overall in the hospital, with the concomitant reduced risk of serious nosocomial (hospital-acquired) infections.

The invention further provides pharmaceutical compositions for treatment of gram-negative bacterial infections, comprising an antibiotic and a BPI protein product in an amount effective to improve the therapeutic effectiveness of the antibiotic. Such compositions optionally comprise pharmaceutically acceptable diluents, adjuvants or carriers. The compositions may be formulated for systemic or topical administration to subjects. In addition, antiseptic compositions comprising BPI protein product and an antibiotic can be used in a variety of *in vitro* uses such as use as a bactericide to decontaminate fluids and surfaces and to sterilize surgical

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and other medical equipment and implantable devices, including prosthetic joints. The invention also provides improved methods of *in vitro* decontamination of fluids and surfaces comprising administering a BPI protein product in combination with an antibiotic.

5 Either the BPI protein product or the antibiotics may be administered systemically or topically. Systemic routes of administration include oral, intravenous, intramuscular or subcutaneous injection, intrathecal, intraperitoneal (e.g. by intraperitoneal lavage), transpulmonary using aerosolized or nebulized drug, or transdermal. Topical routes include administration in the form of salves, ophthalmic
10 drops, ear drops, or irrigation fluids (for, e.g., irrigation of wounds).

 "Concurrent administration," or co-treatment, as used herein includes administration of the agents together, or before or after each other. The BPI protein product and antibiotics may be administered by different routes. For example, the BPI protein product may be administered intravenously while the antibiotics are
15 administered intramuscularly, intravenously, subcutaneously, orally or intraperitoneally. Alternatively, the BPI protein product may be administered intraperitoneally while the antibiotics are administered intraperitoneally or intravenously, or the BPI protein product may be administered in an aerosolized or nebulized form while the antibiotics are administered, e.g., intravenously. The BPI
20 protein product and antibiotics are preferably both administered intravenously. The BPI protein product and antibiotics may be given sequentially in the same intravenous line, after an intermediate flush, or may be given in different intravenous lines. The BPI protein product and antibiotics may be administered simultaneously or sequentially, as long as they are given in a manner sufficient to allow both agents to
25 achieve effective concentrations at the site of infection.

 As used herein, "BPI protein product" includes naturally and recombinantly produced BPI protein: natural, synthetic, and recombinant biologically active polypeptide fragments of BPI protein: biologically active polypeptide variants of BPI protein or fragments thereof, including hybrid fusion proteins and dimers; and
30 biologically active polypeptide analogs of BPI protein or fragments or variants thereof, including cysteine-substituted analogs. The BPI protein products administered according to this invention may be generated and/or isolated by any

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means known in the art. U.S. Patent No. 5,198,541, the disclosure of which is incorporated herein by reference, discloses recombinant genes encoding and methods for expression of BPI proteins including recombinant BPI holoprotein, referred to as rBPI₃₀ and recombinant fragments of BPI. Co-owned, copending U.S. Patent Application Ser. No. 07/885,501 and a continuation-in-part thereof, U.S. Patent Application Ser. No. 08/072,063 filed May 19, 1993 and corresponding PCT Application No. 93/04752 filed May 19, 1993, which are all incorporated herein by reference, disclose novel methods for the purification of recombinant BPI protein products expressed in and secreted from genetically transformed mammalian host cells in culture and discloses how one may produce large quantities of recombinant BPI products suitable for incorporation into stable, homogeneous pharmaceutical preparations.

Biologically active fragments of BPI (BPI fragments) include biologically active molecules that have the same or similar amino acid sequence as a natural human BPI holoprotein, except that the fragment molecule lacks amino-terminal amino acids, internal amino acids, and/or carboxy-terminal amino acids of the holoprotein. Nonlimiting examples of such fragments include a N-terminal fragment of natural human BPI of approximately 25 kD, described in Ooi et al., *J. Exp. Med.*, 174:649 (1991), and the recombinant expression product of DNA encoding N-terminal amino acids from 1 to about 193 or 199 of natural human BPI, described in Gazzano-Santoro et al., *Infect. Immun.* 60:4754-4761 (1992), and referred to as rBPI₁₉₃. In that publication, an expression vector was used as a source of DNA encoding a recombinant expression product (rBPI₁₉₃) having the 31-residue signal sequence and the first 199 amino acids of the N-terminus of the mature human BPI, as set out in Figure 1 of Gray et al., *supra*, except that valine at position 151 is specified by GTG rather than GTC and residue 185 is glutamic acid (specified by GAG) rather than lysine (specified by AAG). Recombinant holoprotein (rBPI) has also been produced having the sequence (SEQ ID NOS: 145 and 146) set out in Figure 1 of Gray et al., *supra*, with the exceptions noted for rBPI₁₉₃ and with the exception that residue 417 is alanine (specified by GCT) rather than valine (specified by GTT). Other examples include dimeric forms of BPI fragments, as described in co-owned and co-pending U.S. Patent Application Serial No. 08/212,132, filed March

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11, 1994, and corresponding PCT Application No. _____ filed March 11, 1995, the disclosures of which are incorporated herein by reference. Preferred dimeric products include dimeric BPI protein products wherein the monomers are amino-terminal BPI fragments having the N-terminal residues from about 1 to 175 to about 1 to 199 of BPI holoprotein. A particularly preferred dimeric product is the dimeric form of the BPI fragment having N-terminal residues 1 through 193, designated rBPI₁₉₃ dimer.

Biologically active variants of BPI (BPI variants) include but are not limited to recombinant hybrid fusion proteins, comprising BPI holoprotein or biologically active fragment thereof and at least a portion of at least one other polypeptide, and dimeric forms of BPI variants. Examples of such hybrid fusion proteins and dimeric forms are described by Theofan et al. in co-owned, copending U.S. Patent Application Serial No. 07/885,911, and a continuation-in-part application thereof, U.S. Patent Application Serial No. 08/064,693 filed May 19, 1993 and corresponding PCT Application No. US93/04754 filed May 19, 1993, which are all incorporated herein by reference and include hybrid fusion proteins comprising, at the amino-terminal end, a BPI protein or a biologically active fragment thereof and, at the carboxy-terminal end, at least one constant domain of an immunoglobulin heavy chain or allelic variant thereof.

Biologically active analogs of BPI (BPI analogs) include but are not limited to BPI protein products wherein one or more amino acid residues have been replaced by a different amino acid. For example, co-owned, copending U.S. Patent Application Ser. No. 08/013,801 filed February 2, 1993 and corresponding PCT Application No. US94/01235 filed February 2, 1994, the disclosures of which are incorporated herein by reference, discloses polypeptide analogs of BPI and BPI fragments wherein a cysteine residue is replaced by a different amino acid. A preferred BPI protein product described by this application is the expression product of DNA encoding from amino acid 1 to approximately 193 or 199 of the N-terminal amino acids of BPI holoprotein, but wherein the cysteine at residue number 132 is substituted with alanine and is designated rBPI₁₉₃Δcys or rBPI₁₉₉. Other examples include dimeric forms of BPI analogs; e.g. co-owned and co-pending U.S. Patent Application Serial No. 08/212,132 filed March 11, 1994, and corresponding PCT

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Application No. _____ filed March 11, 1995, the disclosures of which are incorporated herein by reference.

Other BPI protein products useful according to the methods of the invention are peptides derived from or based on BPI produced by recombinant or synthetic means (BPI-derived peptides), such as those described in co-owned and copending PCT Application No. US94/_____ filed September 15, 1994, which corresponds to U.S. Patent Application Serial No. 08/_____ [Attorney Docket No. 93,1133-H] filed September 15, 1994, and PCT Application No. US94/02465 filed March 11, 1994, which corresponds to U.S. Patent Application Serial No. 08/209,762, filed March 11, 1994, which is a continuation-in-part of U.S. Patent Application Serial No. 08/183,222, filed January 14, 1994, which is a continuation-in-part of U.S. Patent Application Ser. No. 08/093,202 filed July 15, 1993 (for which the corresponding international application is PCT Application No. US94/02401 filed March 11, 1994), which is a continuation-in-part of U.S. Patent Application Ser. No. 08/030,644 filed March 12, 1993, the disclosures of all of which are incorporated herein by reference.

Presently preferred BPI protein products include recombinantly-produced N-terminal fragments of BPI, especially those having a molecular weight of approximately between 21 to 25 kD such as rBPI₂₃ or rBPI₁₁, or dimeric forms of these N-terminal fragments (e.g., rBPI₁₇ dimer). Additionally, preferred BPI protein products include rBPI₃₀ and BPI-derived peptides.

The administration of BPI protein products is preferably accomplished with a pharmaceutical composition comprising a BPI protein product and a pharmaceutically acceptable diluent, adjuvant, or carrier. The BPI protein product may be administered without or in conjunction with known surfactants, other chemotherapeutic agents or additional known anti-microbial agents. A preferred pharmaceutical composition containing BPI protein products (e.g., rBPI₃₀, rBPI₂₃) comprises the BPI protein product at a concentration of 1 mg/ml in citrate buffered saline (5 or 20 mM citrate, 150 mM NaCl, pH 5.0) comprising 0.1% by weight of poloxamer 188 (Pluronic F-68, BASF Wyandotte, Parsippany, NJ) and 0.002% by weight of polysorbate 80 (Tween 80, ICI Americas Inc., Wilmington, DE). Another preferred pharmaceutical composition containing BPI protein products (e.g., rBPI₁₁)

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comprises the BPI protein product at a concentration of 2 mg/ml in 5 mM citrate, 150 mM NaCl, 0.2% poloxamer 188 and 0.002% polysorbate 80. Such preferred combinations are described in co-owned, co-pending PCT Application No. US94/01239 filed February 2, 1994, which corresponds to U.S. Patent Application Ser. No. 08/190,869 filed February 2, 1994 and U.S. Patent Application Ser. No. 08/012,360 filed February 2, 1993, the disclosures of all of which are incorporated herein by reference.

Suitable antibiotics, and therapeutically effective concentrations thereof when administered with BPI protein products, may be determined in *in vivo* models or according to *in vitro* tests, for example, the *in vitro* minimum inhibitory concentration (MIC) and *in vivo* mouse peritonitis or rabbit bacteremia assays taught herein. Suitable antibiotics are antibiotics that act on the bacterial cell wall, cell membrane, protein metabolism or nucleic acid metabolism. These would include antibiotics or combinations of antibiotics from the following classes: β -lactam antibiotics with or without β -lactamase inhibitors, aminoglycosides, tetracyclines, sulfonamides and trimethoprim, vancomycin, macrolides, fluoroquinolones and quinolones, polymyxins, and other antibiotics. Dosage and administration of suitable antibiotics are known in the art, and briefly summarized below.

PENICILLINS

When a BPI protein product is concurrently administered with a penicillin, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 μ g/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily. The penicillin is generally given in doses ranging from 1 μ g/kg to 750 mg/kg daily, preferably not to exceed 24 grams daily for adults (or 600 mg/kg daily for children), and is preferably administered as follows:

Penicillin G is preferably administered parenterally to adults in doses ranging from 600,000 to 1,000,000 units per day. In conventional administration, it is effective largely against gram-positive organisms. For treatment of pneumococcal meningitis, penicillin G is administered in doses of 20-24 million units daily, in divided doses every 2 or 3 hours. For children, the preferred parenteral dose of

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penicillin G is 300,000 to 1,000,000 units per day. One unit of penicillin G contains 0.6 μ g of pure sodium penicillin G (i.e., 1 mg is 1667 units).

Amoxicillin may be administered parenterally to adults in doses ranging from 750 mg to 1.5 grams per day, in 3 equally divided doses. For children, preferred parenteral doses of amoxicillin range from 20 to 40 mg/kg per day in 3 equally divided doses. Amoxicillin is also available in combination with clavulanic acid, a β -lactamase inhibitor. A 250 mg dose of the combination drug amoxicillin/clavulanate will contain 250 mg of amoxicillin and either 125 or 62.5 mg of clavulanic acid. The combination is preferably administered to adults orally in doses of 750 mg per day divided into 3 equal doses every 8 hours, with a preferred dose of 1.5 grams per day for severe infections, given in 3 equally divided doses. In children, the preferred oral dose is 20 to 40 mg/kg per day in 3 equally divided doses.

Ampicillin is preferably administered parenterally to adults in doses of 6 to 12 grams per day for severe infections, in 3 to 4 equally divided doses. In children, the preferred parenteral dose of ampicillin is 50 to 200 mg/kg per day in 3 to 4 equally divided doses. Larger doses of up to 400 mg/kg per day, for children, or 12 grams per day, for adults, may be administered parenterally for treatment of meningitis. Ampicillin is also available in combination with sulbactam, a β -lactamase inhibitor. Each 1.5 gram dose of ampicillin/sulbactam contains 1 gram of ampicillin and 0.5 grams of sulbactam. The combination is preferably administered parenterally to adults in doses of 6 to 12 grams per day divided into 4 equal doses every 6 hours, not to exceed a total of 12 grams per day.

Azlocillin is preferably administered parenterally to adults in doses of 8 to 18 grams per day, given in 4 to 6 equally divided doses.

Carbenicillin is preferably administered parenterally to adults in doses of 30 to 40 grams per day, given by continuous infusion or in 4 to 6 equally divided doses. Daily doses of up to 600 mg/kg have been used to treat children with life-threatening infections.

Mezlocillin is preferably administered to adults parenterally in doses of 100 to 300 mg/kg per day, given in 4 to 6 equally divided doses. The usual dose is 16 to 30 grams per day; for life threatening infections, 350 mg/kg per day may be

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administered, but in doses not to exceed 24 grams per day given in 6 equally divided doses every 4 hours. For children, the preferred parenteral dose of mezlocillin is 150 to 300 mg/kg per day.

5 Nafcillin is preferably administered intravenously to adults in doses of 3 grams per day, given in 6 equally divided doses every 4 hours, with doubled doses for very severe infections. In conventional administration, it is effective largely against gram-positive organisms. In children, the preferred parenteral dose is 20 to 50 mg/kg per day, in 2 equally divided doses every 12 hours. The preferred oral dose for nafcillin ranges from 1 gram per day to 6 grams per day in 4 to 6 divided doses.

10 Oxacillin is preferably administered parenterally to adults in doses of 2 to 12 grams per day, in 4 to 6 equally divided doses. In conventional administration, it is effective largely against gram-positive organisms. In children, oxacillin is preferably administered in doses of 100 to 300 mg/kg per day.

15 Piperacillin is preferably administered parenterally to adults in doses ranging from 100 mg/kg, or 6 grams per day, in 2 to 4 equally divided doses, up to a maximum of 24 grams per day, in 4 to 6 equally divided doses. Higher doses have been used without serious adverse effects.

20 Ticarcillin is preferably administered parenterally to adults in doses ranging from 4 grams per day to 18 grams per day administered in 4 to 6 equally divided doses. The usual dose is 200 to 300 mg/kg per day. For children, the preferred parenteral dose of ticarcillin ranges from 50 mg/kg per day to 300 mg/kg per day, given in 3, 4 or 6 equally divided doses. The combination ticarcillin/clavulanate is preferably administered parenterally to adults in doses of 200 to 300 mg/kg per day (based on ticarcillin content), in 4 to 6 equally divided doses. For adults, the usual dose is 3.1 grams (which contains 3 grams of ticarcillin and 100 mg of clavulanic acid) every 4 to 6 hours. The combination is also available in a dose of 3.2 grams, which contains 3 grams of ticarcillin and 200 mg of clavulanic acid.

30 In general, it is desirable to limit each intramuscular injection of a penicillin or cephalosporin to 2 grams; larger doses should be administered by multiple injections in different large muscle masses.

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CEPHALOSPORINS

When a BPI protein product is concurrently administered with a cephalosporin, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 μ g/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily. The cephalosporin is generally given in doses ranging from 1 μ g/kg to 500 mg/kg daily, preferably not to exceed 16 grams daily, and is preferably administered as follows:

Cefamandole is preferably administered parenterally to adults in doses ranging from 1.5 grams per day, given in 3 equally divided doses every 8 hours, to 12 grams per day for life-threatening infections, given in 6 equally divided doses every 4 hours. In children, cefamandole is preferably administered in doses ranging from 50 to 150 mg/kg per day, in 3 to 6 equally divided doses, not to exceed a total of 12 grams per day.

Cefazolin is preferably administered parenterally to adults in doses of 750 mg per day, given in 3 equally divided doses every 8 hours. In severe, life-threatening infections, it may be administered at doses of 6 grams per day divided into 4 equal doses every 6 hours; in rare instances, up to 12 grams per day have been used. In children, the preferred parenteral dose of cefazolin is 20 to 50 mg/kg per day, divided into 3 or 4 equal doses, with 100 mg/kg per day administered for severe infections.

Cefonicid is preferably administered parenterally to adults in doses ranging from 500 mg once daily, to 2 grams once daily for life-threatening infections. For intramuscular administration, a 2 gram dose should be divided into two 1-gram injections.

Cefoperazone is preferably administered parenterally to adults in doses ranging from 2 grams per day, given in 2 equally divided doses every 12 hours, to 12 grams per day for severe infections, given in 2, 3 or 4 equally divided doses. Doses up to 16 grams per day have been administered without complications.

Cefotetan is preferably administered parenterally to adults in doses of 1 to 4 grams per day, in 2 equally divided doses every 12 hours. Cefotetan may be administered in higher doses for life-threatening infections, not to exceed a total dose of 6 grams per day.

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Cefotaxime is preferably administered parenterally to adults in doses ranging from 1 to 12 grams per day, not to exceed 12 grams per day (2 grams every 4 hours) for life-threatening infections. In children, the parenteral dose of cefotaxime is preferably 50 to 180 mg/kg, divided into 4 to 6 equal doses.

5 Cefoxitin is preferably administered parenterally to adults in doses ranging from 3 to 12 grams per day, given in 3, 4, or 6 equally divided doses. In children, cefoxitin is preferably administered parenterally in doses of 80 to 160 mg/kg per day, given in 4 or 6 equally divided doses, not to exceed a total dose of 12 grams per day.

10 Ceftazidime is preferably administered parenterally to adults in doses ranging from 500 mg per day, given in 2 to 3 equally divided doses (every 8 or 12 hours), up to a maximum of 6 grams per day. In children, ceftazidime is preferably administered intravenously in doses of 30 to 50 mg/kg, to a maximum of 6 grams per day.

15 Cefprozime is preferably administered parenterally to adults in doses ranging from 1 gram per day, given in 2 equally divided doses every 12 hours, to 12 grams per day for life-threatening infections, given in 3 equally divided doses every 8 hours. The usual adult dose is 1 to 2 grams every 8 or 12 hours. For children, the preferred parenteral dose is 50 mg/kg every 6 or 8 hours, for a total daily dose of 200 mg/kg.

20 Ceftriaxone is preferably administered parenterally to adults in doses ranging from 1 to 2 grams per day, given in 2 equally divided doses every 12 hours. It may be given in higher doses, not to exceed a total of 4 grams per day. In children, the preferred parenteral dose of ceftriaxone is 50 to 75 mg/kg per day, not to exceed 2 grams per day. In meningitis, ceftriaxone may be administered in doses of 100 mg/kg per day, not to exceed 4 grams per day.

25 Cefuroxime is preferably administered parenterally to adults in doses ranging from 2.25 to 4.5 grams per day, in 3 equally divided doses every 8 hours. For life-threatening infections, 6 grams per day may be administered in 4 equally divided doses every 6 hours, and for meningitis, 9 grams per day may be administered in 3 equally divided doses every 8 hours. For children, the preferred

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parenteral dose of cefuroxime is 50 to 150 mg/kg per day in 3 to 4 equally divided doses, or 240 mg/kg per day for meningitis.

5 Cephalexin is formulated for oral administration, and is preferably administered orally to adults in doses ranging from 1 to 4 grams per day in 2 to 4 equally divided doses. For children, the preferred dose is 20 to 50 mg/kg per day in divided doses, with doses being doubled for severe infections.

Cephalothin is usually administered parenterally to adults in doses of 8 to 12 grams per day.

OTHER BETA-LACTAMS

10 When a BPI protein product is concurrently administered with an imipenem antibiotic, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 μ g/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily. The imipenem is generally given in doses ranging from 1 μ g/kg to 100 mg/kg daily, and is preferably administered as follows:

15 Imipenem is available in combination with cilastatin, an inhibitor of the renal dipeptidase enzyme that rapidly inactivates imipenem. The combination is preferably administered intramuscularly to adults in doses of 1 to 1.5 grams per day, given in 2 equally divided doses every 12 hours. Intramuscular doses exceeding 1.5
20 grams per day are not recommended. The combination is preferably administered intravenously in doses ranging from 1 to 4 grams per day, in 4 equally divided doses every 6 hours; doses exceeding 50 mg/kg per day, or 4 grams per day, are not recommended.

25 When a BPI protein product is concurrently administered with a monobactam antibiotic, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 μ g/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily. The monobactam is generally given in doses ranging from 1 μ g/kg to 200 mg/kg daily, and is preferably administered as follows:

30 Aztreonam is preferably administered parenterally to adults in doses ranging from 1 gram per day, given in 2 equally divided doses every 12 hours, up

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to a maximum recommended dose of 8 grams per day in cases of life-threatening infection, given in 3 or 4 equally divided doses.

AMINOGLYCOSIDES

5 When a BPI protein product is concurrently administered with an aminoglycoside, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 μ g/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily. The aminoglycoside is generally given in doses ranging from 1 μ g/kg to 20 mg/kg daily, preferably not to exceed 15 mg/kg daily, and is preferably administered as follows:

10 When administering aminoglycosides, it is desirable to measure serum peak and trough concentrations to ensure the adequacy and safety of the dosage. Dosages should generally be adjusted to avoid toxic peak and trough concentrations. Amikacin is preferably administered parenterally to adults and children in doses of 15 mg/kg per day, divided into two or three equal doses every 8 or 12 hours, and not
15 to exceed a total dose of 1.5 grams per day. For uncomplicated infections, a dose of 500 mg amikacin per day, in 2 equally divided doses, may be administered. Dosages should be adjusted to avoid prolonged serum peak concentrations of amikacin above 35 μ g/ml and prolonged trough concentrations greater than 10 μ g/ml.

Gentamicin is preferably administered parenterally to adults in doses
20 of 3 mg/kg per day, in three equally divided doses every 8 hours. For life-threatening infections, up to 5 mg/kg per day in 3 to 4 equally divided doses may be administered, but this dosage should be reduced to 3 mg/kg per day as soon as clinically indicated. For children, gentamicin is preferably administered parenterally in doses of 6 to 7.5 mg/kg per day. Dosages should be adjusted to avoid prolonged
25 serum peak concentrations of gentamicin above 12 μ g/ml and prolonged trough concentrations greater than 2 μ g/ml.

Netilmicin may be administered parenterally to adults in doses ranging from 3 mg/kg per day, in 2 equally divided doses every 12 hours, to 6.5 mg/kg per day for serious systemic infection, in 2 or 3 equally divided doses. In children, the
30 preferred parenteral dose is 5.5 to 8 mg/kg per day, in 2 or 3 equally divided doses. Dosages should be adjusted to avoid prolonged serum peak concentrations of netilmicin above 16 μ g/ml and prolonged serum trough concentrations above 4 μ g/ml.

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Tobramycin is preferably administered parenterally to adults in doses of 3 mg/kg per day, given in three equally divided doses every 8 hours. For life-threatening infections, tobramycin may be administered in doses up to 5 mg/kg per day, in 3 or 4 equally divided doses, but this dosage should be reduced to 3 mg/kg per day as soon as clinically indicated. In children, tobramycin is preferably administered parenterally in doses of 6 to 7.5 mg/kg per day. Prolonged serum concentrations of tobramycin above 12 µg/ml should be avoided, and rising trough levels above 2 µg/ml may indicate tissue accumulation, which may contribute to toxicity.

Concurrent administration of BPI protein product with the aminoglycosides, including amikacin, gentamicin, netilmicin and tobramycin, may permit a lowering of the dose of these toxic antibiotics necessary to achieve a therapeutic effect.

TETRACYCLINES

When a BPI protein product is concurrently administered with a tetracycline, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 µg/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily. The tetracycline is generally given in doses ranging from 1 µg/kg to 50 mg/kg daily, and is preferably administered as follows:

The tetracycline antibiotics are generally administered to adults in doses of 1 to 2 grams per day. An exception is doxycycline, which is preferably administered intravenously to adults in doses of 100 to 200 mg per day, and to children in doses of 2 mg/lb per day. Tetracycline may be administered parenterally to adults in doses of 0.5 to 2 grams per day, in 2 equally divided doses, and to children in doses of 10 to 20 mg/kg per day.

SULFONAMIDES

When a BPI protein product is concurrently administered with a sulfonamide or trimethoprim, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 µg/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily. The sulfonamide or trimethoprim is generally given in doses ranging from 1 µg/kg

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to 150 mg/kg daily, preferably not to exceed a combination dose of 960 mg trimethoprim/4.8g sulfamethoxazole daily, and is preferably administered as follows:

The combination trimethoprim/sulfamethoxazole is available in a formulation containing a 1:5 ratio of trimethoprim and sulfamethoxazole (e.g., 16 mg trimethoprim and 80 mg sulfamethoxazole). The combination is preferably administered intravenously to adults or children in doses of 8 to 10 mg/kg (based on the weight of the trimethoprim component) per day, in 2 to 4 equally divided doses. For *Pneumocystis carinii* infection, the combination can be administered in doses of 20 mg/kg (based on the weight of the trimethoprim component) per day, in 3-4 equally divided doses, to a maximum recommended dose of 960 mg trimethoprim/4.8 g sulfamethoxazole per day. Trimethoprim alone is preferably administered orally to adults in doses of 200 mg per day. Sulfamethoxazole alone is preferably administered orally to adults in doses of 2 to 3 grams per day, and to children orally in doses of 50 to 60 mg/kg per day.

FLUOROQUINOLONES

When a BPI protein product is concurrently administered with a fluoroquinolone or quinolone, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 μ g/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily. The fluoroquinolone or quinolone is generally given in doses ranging from 1 μ g/kg to 50 mg/kg daily, preferably not to exceed 1 gram daily, and is preferably administered as follows:

Norfloxacin is preferably administered orally to adults in doses from 400 to 800 mg daily, divided into two doses every 12 hours. Cinoxacin is preferably administered orally to adults in doses of 1 gram per day, given in 2 or 4 equally divided doses. Ciprofloxacin is preferably administered to adults intravenously in doses from 400 to 800 mg daily, or orally in doses from 500 to 1500 mg daily, divided into two doses every 12 hours. Ofloxacin is preferably administered to adults intravenously in doses from 400 to 800 mg daily, or orally in doses from 400 to 800 mg daily, divided into two doses every 12 hours.

VANCOMYCIN

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When a BPI protein product is concurrently administered with vancomycin, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 μ g/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily. The vancomycin is generally given in doses ranging from 1 mg/kg to 50 mg/kg daily, and is preferably administered parenterally to adults in doses of 2 grams per day, divided into 2 or 4 doses every 6 or 12 hours. In children it is preferably administered in doses of 40 mg/kg, given in 4 equally divided doses every 6 hours. In conventional administration, vancomycin is effective largely against gram-positive organisms.

MACROLIDES

When a BPI protein product is concurrently administered with a macrolide, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 μ g/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily. The macrolide is generally given in doses ranging from 1 μ g/kg to 100 mg/kg daily, and is preferably administered as follows:

Erythromycin is preferably administered intravenously to adults and children in doses of 15 to 20 mg/kg per day, given by continuous infusion or in 4 equally divided doses every 6 hours. Erythromycin can be administered at doses up to 4 grams per day in cases of very severe infection.

Clarithromycin is preferably administered orally to adults in doses of 500 mg to 1 gram daily, in equally divided doses every 12 hours.

Azithromycin is preferably administered orally to adults at a dose of 500 mg on the first day of treatment followed by 250 mg once daily for 4 days, for a total dose of 1.5 grams.

OTHERS

When a BPI protein product is concurrently administered with other antibiotics, for treatment of a gram-negative bacterial infection, the BPI protein product is generally given parenterally in doses ranging from 1 μ g/kg to 100 mg/kg daily, and preferably at doses ranging from 1 mg/kg to 20 mg/kg daily.

Polymyxin B is generally given in doses ranging from 1 unit/kg to 45,000 units/kg daily, and is preferably administered intravenously to adults and

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children in doses of 15,000 to 25,000 units/kg per day, divided into 2 equal doses every 12 hours. It may be administered intramuscularly in doses of 25,000 to 30,000 units/kg per day, although these injections are very painful. Doses of polymyxin B as high as 45,000 units/kg per day have been used in limited clinical studies to treat neonates for *Pseudomonas aeruginosa* sepsis. Polymyxin B is the treatment of choice for *P. aeruginosa* meningitis, and is preferably administered intrathecally to adults and older children in doses of 50,000 units once daily for 3 to 4 days, followed by 50,000 units every other day; in children under two years old, it is administered intrathecally in doses of 20,000 daily for 3 to 4 days, followed by 25,000 units every other day.

Chloramphenicol is preferably administered intravenously to adults in doses of 50 mg/kg per day, in 4 equally divided doses; in exceptional cases, it can be administered in doses up to 100 mg/kg per day. In children, chloramphenicol is preferably administered intravenously in doses of 25 mg/kg per day, although up to 100 mg/kg per day can be administered in cases of severe infection.

Clindamycin is preferably administered parenterally to adults in doses ranging from 600 mg to 4.8 grams per day, given in 2, 3 or 4 equally divided doses. It is recommended that the dose in each intramuscular injection not exceed 600 mg. For children, clindamycin is preferably administered parenterally in doses of 15-40 mg/kg per day, given in 3 or 4 equally divided doses.

Dosages of all antimicrobial agents should be adjusted in patients with renal impairment or hepatic insufficiency, due to the reduced metabolism and/or excretion of the drugs in patients with these conditions. Doses in children should also be reduced, generally according to body weight. Those skilled in the art can readily optimize effective dosages and administration regimens for the BPI protein product and the antibiotics in concurrent administration.

Other aspects and advantages of the present invention will be understood upon consideration of the following illustrative examples. Example 1 addresses the bactericidal effect of gentamicin and BPI when administered to two different strains of *E. coli*. Example 2 addresses the bactericidal effect of gentamicin and BPI for a number of other gram-negative strains. Example 3 addresses the bactericidal effect of polymyxin B and BPI when administered to two different strains

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of *E. coli*. Example 4 addresses the bactericidal effect of polymyxin B and BPI for a number of other gram-negative strains. Example 5 addresses the effect of a BPI protein product with a cephalosporin antibiotic in an *in vivo* mouse peritonitis *E. coli* O111:B4 challenge model. Example 6 also relates to the effect of a BPI protein product with a cephalosporin antibiotic in an *in vivo* mouse peritonitis *E. coli* O111:B4 challenge model. Example 7 relates to the effect of a BPI protein product with a cephalosporin antibiotic in an *in vivo* mouse peritonitis *E. coli* O7:K1 challenge model. Example 8 addresses the effect of a BPI protein product with a cephalosporin antibiotic in an *in vivo* rabbit bacteremia *E. coli* O7:K1 challenge model. Example 9 addresses the effect of a BPI protein product with an aminoglycoside antibiotic in an *in vivo* mouse peritonitis *E. coli* O7:K1 challenge model. Example 10 relates to the effect of a BPI protein product *in vitro* on the antibiotic susceptibility of ceftriaxone-resistant gram-negative organisms. Examples 11-19 address large-scale screening of the antibiotic susceptibility-increasing effect of a BPI protein product on a variety of gram-negative organisms: *Pseudomonas aeruginosa* and other *Pseudomonas* species (Example 11), *E. coli* (Example 12), *Citrobacter* (Example 13), *Klebsiella* (Example 14), *Enterobacter* (Example 15), *Serratia* (Example 16), *Proteus* (Example 17), *Providencia* (Example 18), *Morganella* (Example 19), *Acinetobacter* (Example 20), and *Salmonella* and *Shigella* (Example 21). Example 22 examines the early *in vitro* bactericidal effect of BPI protein product and selected antibiotics on *E. coli* J5, *E. coli* O7:K1, *Enterobacter cloacae* and *Klebsiella pneumoniae*. Example 23 examines the effect of a variety of BPI protein products on several representative organisms, *Acinetobacter anitratus*, *Enterobacter cloacae*, and two strains of *E. coli*. Example 24 relates to the screening of BPI peptides for antibacterial activity against *E. coli*. Example 25 addresses the effect on *E. coli* O111:B4 of concurrent administration of BPI protein product with tetracycline or gentamicin.

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Example 1**SYNERGISTIC BACTERICIDAL EFFECTS OF GENTAMICIN AND BPI:
ADMINISTRATION TO *E. COLI* AND *E. COLI* 0111:B4
GRAM-NEGATIVE ORGANISMS**

5 In this example, a micro dilution plate minimum inhibitory concentration (MIC) assay was conducted to determine the sensitivity of *E. coli* organisms to the bactericidal effects of BPI protein products concurrently administered with the antibiotic gentamicin. The assays were conducted against the BPI sensitive organism *E. coli* J5 (an Rc rough mutant of *E. coli* 0111:B4), and a BPI
10 resistant organism *E. coli* 0111:B4.

 Specifically, organisms were grown overnight on blood agar plates at 37°C in air, single colonies were then sub-cultured in 100 mL of nutrient broth No. 2 (Oxoid CM67) and incubated with gentle agitation on an orbital shaker for 5 1/4 hours until in log phase. Fifty mL of the bacterial suspension was then spun down
15 in a Denley BR401 bench centrifuge at 4000 rpm for 15 minutes and the pellet was resuspended and washed twice using sterile normal saline. Bacteria were then resuspended in saline such that a 1:10 dilution had an optical density of 0.9 (+/-0.01) at 325 nm (corresponding to approximately 4×10^8 cells per mL) and diluted to give a final concentration of 4×10^6 cells per mL in "BPI media" (described below).

20 All assays and dilutions of BPI protein products and gentamicin were performed using "BPI media" consisting of 50% peptone water (Oxoid L37, Lot: 25851279) with 0.1 M MOPS (Sigma M-1254) buffered to pH 6.00 with sodium hydroxide. This provides a nutritive media with a low protein and divalent cation concentration adjusted to a pH which is not inhibitory to bacterial growth and allows
25 for readily measurable (though not optimal) BPI activity.

 rBPI₂₂ and gentamicin (Sigma G-1264, Lot: 91H00325) were diluted in BPI media such that 100 µL of diluted BPI, 50 µL of diluted gentamicin and 50 µL of bacterial suspension in the final volume of 200 µL per well, gave concentrations in serial dilutions from 1000 nM (25 µg per mL) BPI and 32 µg per
30 mL of gentamicin with a fixed concentration of 10^6 cells/mL. Checkerboards were then constructed in round bottomed 96 well microtitre plates (Greiner No. 650180) and incubated with non-sealing lids at 37 C in air for 18 hours. Plates were then read

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by eye and in an automatic plate reader (Titretek Multiscan plus) at 580 nm with visible growth corresponding to an optical density of approximately 0.1 (see Table 1 for *E. coli* J5, and Table 3 for *E. coli* O111:B4). Viable counts were made from the wells adjacent to the cutoff of visible growth for the two *E. coli* plates (see Table 2 for *E. coli* J5, and Table 4 for *E. coli* O111:B4) by dilution of 10 μ L in 990 μ L of Nutrient Broth No. 2 and spread plates were prepared with 25 μ L on blood agar. The inhibitory activity of BPI protein product with antibiotic was evaluated by the method of Eliopoulos and Moellering In *Antibiotics in Laboratory Medicine*, 3rd ed. (Lorian, V., Ed.) pp. 432-492, Williams and Wilkins, Baltimore MD (1991), wherein a fractional inhibitory concentration index (FIC) of less than 0.5 was scored as synergy, 1 was scored as additive and greater than 1 but less than 2 was scored as indifferent.

A positive synergistic interaction was demonstrated between rBPI₂₂ and gentamicin using the BPI sensitive *E. coli* J5 with gentamicin at a concentration of 0.25 μ g per mL, reducing the MIC of BPI by approximately eight-fold from 500 nM (12.5 μ g per mL) to 62.5 nM (1.56 μ g per mL) with similar reductions of the minimum bactericidal concentration (MBC).

Visible Growth (bold line) of *E. coli* J5 and ODS (580 nm)

BPI (nM):	Gentamicin (μg per ml)											
	32	16	8	4	2	1	0.5	0.25	1.25	0.6	0.3	0
1000	.071	.067	.069	.067	.068	.071	.073	.072	.065	.068	.062	.065
500	.055	.050	.052	.055	.053	.056	.054	.059	.106	.054	.056	.054
250	.056	.055	.053	.057	.058	.057	.055	.060	.055	.054	.194	.201
125	.058	.053	.051	.053	.055	.062	.064	.061	.060	.215	.258	.236
62.5	.052	.053	.056	.054	.054	.053	.062	.056	.172	.196	.262	.261
31	.057	.058	.052	.055	.056	.058	.063	.135	.206	.251	.259	.286
16	.057	.059	.057	.058	.059	.062	.081	.148	.240	.262	.297	.327
0	.060	.053	.060	.057	.054	.058	.071	.151	.252	.285	.330	.326

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TABLE 2
Visible Growth (bold line) of *E. coli* J5 and Viable Counts

BPI (nM):	Gentamicin (μg per ml)											
	32	16	8	4	2	1	0.5	0.25	.125	.06	.03	0
1000	NC	NC	NC	NC	NC	NC	NC	0	0	0	0	0
500	NC	NC	NC	NC	NC	NC	0	0	147	2	0	0
250	NC	NC	NC	NC	NC	0	0	0	0	0	0	NC
125	NC	NC	NC	NC	0	0	0	0	0	++	NC	NC
62.5	NC	NC	NC	NC	0	0	121	1	++	NC	NC	NC
31	NC	NC	NC	NC	0	0	555	++	NC	NC	NC	NC
16	NC	NC	NC	NC	0	0	438	438	NC	NC	NC	NC
0	NC	NC	NC	NC	0	0	267	++	NC	NC	NC	NC

Counts expressed as counts per ml, NC=not counted, ++=too numerous to count.

TABLE 3

Visible Growth (bold line) of *E. coli* 0111:B4 and ODs (580 nm)

BPI (nM):	Gentamicin (μg per ml)											
	32	16	8	4	2	1	0.5	0.25	.125	.06	.03	0
1000	.067	.064	.057	.057	.055	.067	.062	.191	.275	.328	.308	.370
500	.054	.051	.049	.052	.050	.050	.048	.179	.245	.282	.276	.342
250	.054	.053	.053	.053	.058	.061	.054	.161	.245	.282	.269	.345
125	.048	.051	.049	.047	.049	.050	.109	.198	.239	.267	.261	.328
62.5	.062	.058	.054	.051	.053	.052	.059	.176	.252	.283	.287	.353
31	.059	.054	.054	.052	.051	.051	.054	.192	.250	.287	.278	.357
16	.061	.054	.058	.063	.052	.054	.048	.158	.261	.285	.284	.349
0	.054	.058	.055	.049	.050	.053	.079	.155	.280	.309	.322	.374

TABLE 4

Visible Growth (bold line) of *E. coli* 0111:B4 and viable counts

BPI (nM):	Gentamicin (μ g per ml)									
	32	16	8	4	2	1	0.5	0.25	0.06	0
1000	NC	NC	NC	NC	0	0	6	++	NC	NC
500	NC	NC	NC	NC	0	0	0	+++	NC	NC
250	NC	NC	NC	NC	0	0	0	+++	NC	NC
125	NC	NC	NC	NC	0	0	0	+++	NC	NC
62.5	NC	NC	NC	NC	0	0	0	+++	NC	NC
31	NC	NC	NC	NC	0	0	0	+++	NC	NC
16	NC	NC	NC	NC	0	0	1	+++	NC	NC
0	NC	NC	NC	NC	0	0	0	+++	NC	NC
	NC	NC	NC	NC	0	0	287	+++	NC	NC

Counts expressed as count per ml, NC=not counted, ++=100 numerous to count.

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Example 2**SYNERGISTIC BACTERICIDAL EFFECTS OF GENTAMICIN AND BPI:
ADMINISTRATION TO ADDITIONAL GRAM-NEGATIVE ORGANISMS**

In this example, micro dilution plate MIC assays were conducted
5 according to the method of Example 1 to determine the sensitivity of a variety of
gram-negative organisms to the cytotoxic effects of BPI protein products concurrently
administered with gentamicin antibiotic. The results of those assays are shown in
Table 5 below. Positive synergistic bactericidal interactions were observed against
E. coli JS, *E. coli* 01:K1, *E. coli* (S2252), *K. oxytoca*, *E. tarda*, *Salmonella*
10 *typhimurium* (S2136) and *Salmonella arizonae*. Possible synergistic bactericidal
interactions (indicated as "additive (+)") were observed against *E. cloacae* (10005),
E. gergoviae (11434), *P. aeruginosa* (10332) and *P. aeruginosa* (10662). The
concurrent administration of gentamicin and rBPI₂₃ had only additive, indifferent or
indeterminate effects against the other tested gram-negative bacteria.

TABLE 5
MIC Values and Results of Checkerboard Susceptibility Testing With BPI and Gentamicin

Organism (NCTC)	MIC of Single Agent BPI ₅₀ (nM)	MIC of Single Agent Gentamicin (μg/ml)	Lowest MIC of the Agents Together rBPI ₅₀ /Gentamicin	Interpretation (Gentamicin)
<i>E. coli</i> J5	250	1.0	62.5/0.125	synergy
<i>E. coli</i> O1:K1	>2000	2.0	31/0.5	synergy
<i>E. coli</i> (S2252) NCTC 10418	500	0.5	250/0.25	additive/synergy
<i>E. coli</i> O111:B4	>2000	1.0	>2000/1.0	indifferent
<i>E. coli</i> (S2216)	>500	1.0	>500/1.0	indifferent
<i>E. coli</i> H262	>500	indifferent
<i>E. dispar</i> (S2162)	>500	indifferent
<i>E. alcalescens</i> (S2196)	>2000	indifferent
<i>K. oxytoca</i>	>2000	1.0	1000/0.25	synergy
<i>K. pneumoniae</i>	>2000	0.5	>2000/0.5	indifferent
<i>E. cloacae</i> (10005)	>2000	1.0	1000/0.5	additive (+)
<i>E. gergoviae</i> (11434)	>2000	2.0	2000/0.5	additive (+)
<i>S. marcescens</i> (10211)	>2000	4.0	>2000/4.0	indifferent
<i>P. reingert</i> (S2253)	>500	2.0	1000/1.0	indifferent
<i>P. vulgaris</i> (4175)	500	0.25	250/0.125	additive
<i>P. morganii</i> (S2161)	>2000	2.0	2000/2.0	indifferent
<i>P. aeruginosa</i> (10332)	>2000	0.5	31/0.25	additive (+)
<i>P. aeruginosa</i> (840P)	>2000	1.0	>2000/1.0	indifferent

TABLE 5
MIC Values and Results of Checkerboard Susceptibility Testing With BPI and Gentamicin

Organism (NCTC)	MIC of Single Agent BPI ₁₃ (nM)	MIC of Single Agent Gentamicin (µg/ml)	Lowest MIC of the Agents Together rBPI ₁₃ /Gentamicin	Interpretation (Gentamicin)
<i>P. aeruginosa</i> (10662)	> 2000	0.25	2000/0.125	additive (+)
<i>P. aeruginosa</i> (U600)	> 2000	0.5	62.5/0.25	indifferent
<i>K. aerogenes</i> NCTC 9496	> 2000	1.0	2000/0.5	indifferent
<i>Shigella dysenteriae</i> NCTC 4837	> 2000	1.0	> 2000/1.0	indifferent
<i>E. tarda</i>	500	1.0	31/0.125	synergy
<i>K. rhinoscleromatis</i>	> 500	1.0	> 500/1.0	indifferent
<i>Salmonella choleraesuis</i>	> 2000	---	---	indifferent
<i>Salmonella typhimurium</i> (S2185)	> 500	0.25	> 500/0.25	indifferent
<i>Salmonella typhimurium</i> (S2136)	> 2000	> 4.0	31/4.0	synergy
<i>Salmonella arizonae</i>	> 500	1.0	---	synergy
<i>E. aerogenes</i> (S2164)	> 2000	1.0	62.5/1.0	indifferent
<i>C. freundii</i>	> 2000	---	---	indifferent

(+) possibility of synergy but FIC in calculable because rBPI₁₃ MIC outside tested concentration range.

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Example 3**SYNERGISTIC BACTERICIDAL EFFECTS OF POLYMYXIN B AND BPI:
ADMINISTRATION TO *E. COLI* J5 AND
E. COLI 0111:B4 GRAM-NEGATIVE ORGANISMS**

5 In this example, micro dilution plate MIC assays were conducted according to the method of Example 1 to determine the sensitivity of *E. coli* J5 and *E. coli* 0111:B4 to the cytotoxic effects of BPI protein products concurrently administered with the antibiotic polymyxin B. Polymyxin B solution with an activity of 10240 units/mL was prepared by preparing a solution of 1.595 mg of polymyxin
10 B sulfate stock powder per mL, and diluting it in sterile water for injection as 20 µg in 12.54 mL.

 A positive synergistic interaction was demonstrated between rBPI₂₂ and polymyxin B using *E. coli* 0111:B4 (see Tables 8 and 9) but the administration of polymyxin B with BPI did not have synergistic bactericidal effects with BPI when
15 applied to *E. coli* J5 as illustrated by the results shown in Tables 6 and 7.

TABLE 6

Visible Growth (bold line) of *E. coli* J5 and ODs (580 nm) Polymyxin B ($\mu\text{g/ml}$)

BPI (nM):	5	2.5	1.25	.6	.3	.15	.075	.075	0
2000	.071	.006	.001	.001	.002	.001	.011	.002	.006
1000	.069	.003	.002	.004	.003	.001	.001	.009	.007
500	.055	.000	.018	.001	.001	.007	.012	.012	.006
250	.049	.000	.001	.001	.012	.026	.025	.013	.006
125	.047	.002	.095	.123	.114	.107	.130	.093	.157
62.5	.047	.002	.047	.131	.144	.164	.141	.124	.185
31	.044	.001	.094	.160	.163	.184	.183	.168	.238
0	.046	.002	.128	.230	.254	.282	.270	.245	.307

TABLE 7

Visible Growth (bold line) of *E. coli* J5 and Viable Counts Polymyxin B ($\mu\text{g/ml}$)

BPI (nM):	5	2.5	1.25	.6	.3	.15	.075	.075	0
2000	NC	NC	0	0	0	0	0	0	0
1000	NC	NC	0	5	1	0	0	0	0
500	NC	NC	12	15	++	++	264	379	0
250	NC	81	++	++	+++	+++	+++	+++	496
125	0	0	NC	NC	NC	NC	NC	NC	NC
62.5	0	0	NC	NC	NC	NC	NC	NC	NC
31	0	0	NC	NC	NC	NC	NC	NC	NC
0	0	0	NC	NC	NC	NC	NC	NC	NC

Counts expressed as counts per ml, NC=not counted, ++=too numerous to count.

TABLE 8

Visible Growth (bold line) of *E. coli* 0111:B4 and ODS (580 nM)

BPI (nM):	Polymyxin B (μ g/ml)									
	10	5	2.5	1.25	.6	.3	.15	.075	0	
2000	0	.015	.009	.01	.011	.011	.295	.379	.360	
1000	0	.003	.035	.109	.148	.217	.261	.305	.286	
500	0	.006	.008	.145	.166	.280	.238	.292	.282	
250	0	.006	.012	.146	.216	.265	.254	.294	.260	
125	0	.016	.101	.193	.228	.269	.233	.298	.269	
62.5	0	.050	.012	.212	.201	.277	.246	.288	.281	
31	0	.002	.054	.245	.223	.260	.234	.291	.271	
0	0	.001	.144	.267	.265	.295	.272	.309	.290	

TABLE 9

Visible Growth (bold line) of *E. coli* 0111:B4 and Viable Counts

BPI (nM):	Polymyxin B (μg/ml)									
	10	5	2.5	1.25	.6	.3	.15	.075	0	
2000	NC	0	0	0	0	1	+++	NC	NC	
1000	0	0	NC	++	+++	+++	NC	NC	NC	
500	0	0	0	+++	NC	NC	NC	NC	NC	
250	0	0	89	+++	NC	NC	NC	NC	NC	
125	0	7	+++	+++	NC	NC	NC	NC	NC	
62.5	0	80	5	+++	NC	NC	NC	NC	NC	
31	0	0	+++	NC	NC	NC	NC	NC	NC	
0	0	0	+++	NC	NC	NC	NC	NC	NC	

Counts expressed as counts per ml, NC=not counted, ++=too numerous to count.

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Example 4

**SYNERGISTIC BACTERICIDAL EFFECTS OF POLYMYXIN B AND BPI:
ADMINISTRATION TO ADDITIONAL GRAM-NEGATIVE ORGANISMS**

In this example, micro dilution MIC assays were conducted according
5 to the method of Example 1 to determine the sensitivity of a variety of gram-negative
organisms to the cytotoxic effects of BPI protein products concurrently administered
with polymyxin B antibiotic. The results of those assays are shown in Table 10
below.

These assay results show additive or synergistic effects with the use of
10 polymyxin B at concentrations of 0.3 $\mu\text{g/mL}$, a level of which is 10-20 times lower
than when that antibiotic is used conventionally.

TABLE 10

MIC Values and Results of Checkerboard Susceptibility Testing With BPI and Polymyxin B (PB)				
Organism (NCTC)	MIC of Single Agent BPI ₁₀ (nM)	MIC of Single Agent PB (μg/ml)	Lowest MIC of the Agents Together rBPI ₁₀ /PB	Interpretation (Polymyxin B)
<i>E. coli</i> J5	250	2.5	250/2.5	indifferent
<i>E. coli</i> (S2252) NCTC 10418	500	5.0	250/2.5	indifferent
<i>E. coli</i> 0111:B4	> 2000	5.0	250/2.5	synergy
<i>E. cloacae</i> (10005)	> 500	0.5	500/0.5	indifferent
<i>E. gergoviae</i> (11434)	> 2000	5.0	125/2.5	additive (+)
<i>S. marcescens</i> (10211)	> 2000	> 10.0	> 2000/10.0	indifferent
<i>P. reuteri</i> (S2253)	> 2000	2.5	125/1.25	synergy
<i>P. aeruginosa</i> (10332)	> 2000	1.25	250/0.6	synergy
<i>P. aeruginosa</i> (840P)	> 2000	1.25	1000/0.6	synergy
<i>P. aeruginosa</i> (10662)	> 2000	1.25	125/0.6	synergy
<i>P. aeruginosa</i> (U600)	> 2000	1.25	2000/0.3	synergy
<i>E. aerogenes</i> (S2164)	> 2000	> 10.0	> 2000/10.0	indifferent

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Example 5

SYNERGISTIC EFFECTS OF CEFAMANDOLE AND
BPI PROTEIN PRODUCT *IN VIVO* IN MICE CHALLENGED
INTRAPERITONEALLY WITH LIVE *E. COLI* 0111:B4 BACTERIA:
EFFECT ON SURVIVAL

In this example, the protective effect of cefamandole nafate antibiotic (MANDOL®, Lilly) a semisynthetic broad-spectrum cephalosporin antibiotic with and without a BPI protein product was evaluated by means of challenging ICR mice with an LD₅₀₋₁₀₀ dose level of live *E. coli* 0111:B4 bacteria, a strain that is not susceptible to the bactericidal/growth inhibitory effects of BPI protein product. Specifically, four groups of 15 ICR mice were treated such that each ICR mouse received an injection of bacteria (1.8×10^8 CFU/mouse) intraperitoneally; an intraperitoneal injection of cefamandole nafate (MANDOL®, 100 mg/kg) or saline; and then an intraperitoneal injection of rBPI₂₁ (500 µg/mouse) or BPI buffer. Survival of the mice was then evaluated over a period of 7 days with the results illustrated in Figure 1.

Concurrent administration of BPI protein product with cefamandole, or administration of cefamandole alone, showed significant protection in mice challenged with *E. coli* 0111:B4 when compared to the buffer control ($p \leq 0.001$ and $p \leq 0.05$ respectively). The protective effect of the BPI protein product when administered without cefamandole was not evident. When compared to treatment with cefamandole alone, concurrent administration of BPI protein product with cefamandole showed improved protection at $p \leq 0.1$. These results indicate that the concurrent administration of cefamandole and a BPI protein product has synergistic therapeutic effects against *E. coli* 0111:B4.

Example 6

SYNERGISTIC EFFECTS OF CEFAMANDOLE AND
BPI PROTEIN PRODUCT *IN VIVO* IN MICE CHALLENGED
INTRAPERITONEALLY WITH LIVE *E. COLI* 0111:B4 BACTERIA:
EFFECT ON SURVIVAL AND ON BACTERIAL CLEARANCE
FROM BLOOD AND PERITONEAL LAVAGE FLUID

The protective effects of a cephalosporin antibiotic and a BPI protein product were evaluated in mice challenged intraperitoneally with *E. coli*

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O111:B4, a strain that is resistant to the bactericidal effects of BPI protein product. The assay was conducted using the following procedure. Male ICR mice (Simonsen Laboratories, Gilroy California), 5-7 weeks old, were housed under controlled climate and dark/light cycles and were allowed free access to food and water. Mice received an intraperitoneal injection of 0.5 ml of bacteria in doses near an LD₅₀ (2×10^9 CFU/mouse). Immediately after bacterial challenge the animals received an intraperitoneal injection of (1) vehicle only, (2) 500 µg/mouse rBPI₂₁ and vehicle, (3) 100 mg/kg cefamandole nafate (Mandol[®] in phosphate buffered saline; Eli Lilly, Indianapolis, IN) and vehicle, or (4) 500 µg rBPI₂₁ and 100 mg/kg cefamandole.

Survival of the four groups was monitored for 7 days. Survival data was statistically analyzed using the Chi-square test. In a separate experiment using a bacterial challenge of 2.5×10^9 CFU *E. coli* O111:B4, blood and peritoneal lavage fluid were collected for culture at different time points following bacterial challenge. Blood was obtained from the retro-orbital sinus. At least 1 ml of peritoneal lavage fluid was obtained after intraperitoneal injection of 3 ml of phosphate buffered saline. Bacterial counts (expressed as CFU/ml) were determined by inoculating trypticase soy agar plates with 10-fold dilutions of blood or peritoneal lavage samples, incubating the plates overnight at 37°C, and counting the colonies. Statistical comparisons of this data were performed with the analysis of variance.

Figure 2 shows effects on survival of cefamandole (100 mg/kg), rBPI₂₁ (500 µg) or the concurrent administration of the two agents. The BPI protein product alone had no effect, while cefamandole treatment alone increased survival, although not significantly. In contrast, the concurrent administration of rBPI₂₁ with cefamandole resulted in a significant increase in survival ($p < 0.05$) above that achieved by either treatment alone. Figures 3 and 4 show that rBPI₂₁ alone failed to reduce counts in either peritoneal lavage fluid or blood after challenge with bacteria, while cefamandole treatment alone significantly reduced counts in both ($p < 0.01$ vs. vehicle). However, the concurrent administration of rBPI₂₁ and cefamandole reduced bacterial counts in the peritoneal lavage fluid by more than two orders of magnitude below that achieved by cefamandole alone

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($p < 0.01$ of concurrent administration vs. cefamandole only) at the 2 and 6 hour time points, and entirely eliminated counts after 24 hours. Blood of mice that received concurrent administration of rBPI₂₁ and cefamandole was completely free of bacteria at all time points.

5 Thus, the concurrent administration of a BPI protein product, rBPI₂₁, with a suboptimal dose of a cephalosporin antibiotic, cefamandole, resulted in a superior therapeutic effect. The data indicates that BPI protein products and cephalosporin antibiotics produce a synergistic therapeutic effect. Since cefamandole alone reduced counts by approximately two orders of magnitude
10 compared to vehicle-treated animals, another experiment was conducted to determine if rBPI₂₁ alone reduced bacterial counts when the inoculum was reduced to 10^7 CFU. Data from this experiment showed that rBPI₂₁ (500 μ g) did not significantly reduce bacterial counts in blood or peritoneal lavage fluid after a challenge of 10^7 CFU. This suggests that an antibiotic-mediated reduction in the
15 magnitude of the bacterial count cannot, in itself, explain the protection associated with concurrent administration of BPI protein product and antibiotic.

Example 7

20 SYNERGISTIC EFFECTS OF CEFAMANDOLE AND BPI PROTEIN PRODUCT *IN VIVO* IN MICE CHALLENGED INTRAPERITONEALLY WITH LIVE *E. COLI* O7:K1 BACTERIA: EFFECT ON SURVIVAL

The protective effects of a cephalosporin antibiotic and a BPI
protein product were evaluated in mice challenged intraperitoneally with *E. coli*
25 O7:K1 (ATCC Accession No. 23503), a strain that is susceptible to the bactericidal effects of BPI protein product. The general procedure described above in Example 5 was followed. Four groups of 20 mice were challenged intraperitoneally with 2×10^7 *E. coli* O7:K1 bacteria and then treated with (1)
vehicle, (2) 50 μ g rBPI₂₁ only, (3) 20 mg/kg cefamandole only, or (4) both 50 μ g
30 rBPI₂₁ and 20 mg/kg cefamandole. Survival of the mice was followed over a period of 7 days; results of two trials are displayed in Figures 5A and 5B.

In one trial, rBPI₂₁ alone protected 11% of the survivors compared to vehicle controls. cefamandole protected 47% of the survivors compared to

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vehicle controls ($p < 0.05$ vs. vehicle), and the concurrent administration of rBPI₂₁ and cefamandole protected 100% of the survivors compared to vehicle controls ($p < 0.001$ vs. vehicle, $p < 0.01$ vs. cefamandole alone). In the second trial, rBPI₂₁ alone protected 0% of the survivors compared to vehicle, cefamandole protected 12% compared to vehicle, and the concurrent administration of rBPI₂₁ and cefamandole protected 59% compared to vehicle ($p < 0.01$ vs. vehicle, $p < 0.05$ vs. cefamandole alone). In both trials, the increase in survival associated with the concurrent administration of rBPI₂₁ and cefamandole was greater than the sum of the increases in survival due to the individual therapies. Thus, there appears to be synergy (a greater than additive effect) between cefamandole and BPI protein product in this model.

Example 8

SYNERGISTIC EFFECTS OF CEFAMANDOLE AND BPI PROTEIN PRODUCT *IN VIVO* IN RABBITS CHALLENGED INTRAVENOUSLY WITH LIVE *E. COLI* O7:K1 BACTERIA: EFFECT ON BACTERIAL CLEARANCE AND ON CARDIOVASCULAR, RESPIRATORY AND METABOLIC PARAMETERS

Adult male New Zealand White rabbits (Charles River Laboratories, St. Constant, Canada) weighing between 1.8 and 2.3 kilograms were fasted for 24 hours before the experiment. Each rabbit was anesthetized with an intramuscular injection of 80/4 mg/kg Ketamine/xylazine. The left femoral artery was catheterized for blood pressure determinations and blood sample collection. A catheter was placed adjacent the right atrium via the right jugular vein and a thermistor-tipped catheter was placed in the aortic arch via the right carotid artery. The rabbits were allowed to stabilize for 90-120 minutes following catheterization to normalize hemodynamic and blood gas parameters.

The rabbits were divided into four treatment groups with 4 animals per group: (1) vehicle alone, (2) cefamandole and vehicle, (3) rBPI₂₁ and vehicle, and (4) cefamandole and rBPI₂₁. The rabbits were administered cefamandole (Mandol[®]; Eli Lilly, Indianapolis, IN) or vehicle intravenously 5 minutes before the start of the bacterial infusion (considered to be $T=0$). At $T=0$, 2×10^{10} CFU/rabbit of *E. coli* O7:K1 was infused intravenously over 10 minutes into the

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ear vein. Simultaneously (at T=0), 10 mg/kg rBPI₂₁ or vehicle was infused over 10 minutes via the right jugular catheter. After the ten minute infusion, rBPI₂₁ was slowly infused at 10 mg/kg/hr for 2 hours (resulting in a total dose of 30 mg/kg rBPI₂₁).

- 5 Arterial blood samples for determination of bacterial counts and endotoxin levels were collected at the end of the 10 minute bacterial infusion and at 30 minutes, 1, 2, 3 and 4 hours. The whole blood was 10-fold serially diluted in sterile PBS and aliquots were plated onto tryptic soy agar plates, incubated at 37°C overnight, and the plates were counted for colony forming units (CFU). The
10 results were expressed as CFU/ml blood and percent bacteria dose per ml blood. The remaining portion of the blood was centrifuged, the plasma removed and passed through a 0.2 micron Whatman syringe filter to remove the bacteria. The endotoxin levels were determined using a modified limulus amoebocyte lysate assay (Pyrochrome LAL Assay, Associates of Cape Cod, Woods Hole, MA).
15 These results were expressed as ng LPS per ml plasma.

- Cardiovascular, respiratory and metabolic parameters were measured every 30 minutes. Mean arterial blood pressure (MABP) and heart rate were monitored continuously throughout the experiments and displayed on a cardiac output computer (Columbus Instruments Cardiomax II) or on a chart
20 recorder. Heart rate was derived from the arterial pressure wave. Cardiac output was determined in duplicate with the thermodilution technique: Changes in blood temperature resulting from injection of 900 µl of room temperature PBS were recorded with the thermistor-tipped catheter in the aortic arch. The cardiac output computer then generated thermodilution curves that were visualized on the chart
25 recorder, and derived cardiac output from the temperature-time curves. Cardiac index (CI) was then calculated as cardiac output per kg body weight. In addition, total peripheral resistance (TPR) was determined by dividing blood pressure by cardiac output.

- Blood samples for blood gas determinations were drawn from the
30 femoral artery catheter every 30 minutes throughout the study. Blood gases were measured with a Ciba-Corning Blood Gas System, Model 278 (Ciba-Corning Diagnostics Corp., Medfield, MA). The blood gas system directly measures blood

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pH, partial pressure of pCO_2 , and partial pressure of pO_2 . Other parameters including the alveolar-arterial oxygen gradient, arterial oxygen content, estimated oxygen saturation, standard bicarbonate, and *in vivo* base excess were calculated using the formulas provided by Ciba-Corning Diagnostic Corp. Plasma levels of glucose and lactate were determined using a Glucose/L-lactate Analyzer (2300 STAT, YSI, Yellow Springs, OH).

Survival data is shown in Figure 6. Two of the four animals (50%) treated with vehicle alone died before the end of the experiment. None of the animals concurrently treated with $rBPI_{11}$ and cefamandole died. Bacterial counts in blood, expressed as CFU/ml or percent bacteria dose/ml, are shown in Figures 7 and 8, respectively. In Figure 7, the squares represent treatment with vehicle alone, the diamonds represent cefamandole alone, the circles represent $rBPI_{11}$ alone, and the triangles represent the concurrent administration of $rBPI_{11}$ and cefamandole. In Figure 8, the bar with horizontal hatching indicates treatment with cefamandole alone, the hollow bar indicates the concurrent administration of $rBPI_{11}$ and cefamandole, the bar with vertical hatching indicates $rBPI_{11}$ alone, and the solid bar indicates buffer alone. The group concurrently treated with $rBPI_{11}$ and cefamandole demonstrated a higher clearance of bacteria compared to the groups treated with either $rBPI_{11}$ alone or cefamandole alone. There appears to be a synergistic effect at 30 and 60 minutes; at 30 minutes, the concurrent administration of $rBPI_{11}$ and cefamandole resulted in a higher percentage clearance of bacteria than the sum of the separate treatments.

Endotoxin levels are displayed in Figure 9. The open diamonds indicate treatment with vehicle alone, filled diamonds indicate $rBPI_{11}$ alone, the filled squares indicate cefamandole alone, and the open squares indicate concurrent administration of $rBPI_{11}$ and cefamandole. Animals administered cefamandole alone have a much higher LPS level than animals treated with vehicle alone, due to release of LPS as cefamandole kills bacteria. The concurrent administration of $rBPI_{11}$ and cefamandole produced a dramatic decrease in LPS levels compared to cefamandole therapy alone.

Cardiovascular/pulmonary parameters (MABP, CI, TPR, arterial oxygen tension, alveolar-arterial O_2 gradient, respiration rate, and arterial blood

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pH) are shown in Figures 10-16, respectively. These figures only display results for the groups treated with cefamandole alone (indicated by open squares) and concurrent treatment with rBPI₂₁ and cefamandole (indicated by filled squares). Single stars indicate that the concurrent administration of both agents provided statistically significant ($p < 0.05$) improvement over antibiotic alone, while two stars indicate $p < 0.01$. Cefamandole alone or rBPI₂₁ alone failed to protect the animals; cardiovascular and respiratory dysfunction began during the bacterial infusion and the animals were in circulatory shock by the end of the infusion. Cardiovascular shock lasted for the rest of the experiment. Arterial blood pH began to decrease at 60 minutes in the group treated with cefamandole alone, and was at its lowest level by the end of the experiment. In contrast, concurrent administration of rBPI₂₁ and cefamandole preserved cardiopulmonary function and prevented septic shock. Thus, the concurrent administration of BPI protein product with antibiotic protected the animals against the lethal effect of bacteremia and preserved cardiopulmonary function when antibiotic alone failed to do so.

Example 9

SYNERGISTIC EFFECTS OF GENTAMICIN AND BPI PROTEIN PRODUCT *IN VIVO* IN MICE CHALLENGED INTRAPERITONEALLY WITH LIVE *E. COLI* O7:K1 BACTERIA: EFFECT ON SURVIVAL

The synergistic effects of an aminoglycoside antibiotic and a BPI protein product were evaluated in mice challenged intraperitoneally with *E. coli* O7:K1 (ATCC Accession No. 23503), a smooth encapsulated strain that is susceptible to the bactericidal effects of BPI protein product. The general procedure described above in Example 4 was followed. Six groups of 20 mice were challenged intraperitoneally with 2×10^7 *E. coli* O7:K1 bacteria and treated immediately with (1) vehicle, (2) 0.03 mg/kg gentamicin, (3) 0.1 mg/kg of gentamicin, (4) 50 μ g of rBPI₂₁, (5) 0.03 mg/kg gentamicin followed by 50 μ g of rBPI₂₁, or (6) 0.1 mg/kg of gentamicin followed by 50 μ g of rBPI₂₁. Survival was followed over 7 days. Results are shown in Figure 17. Neither antibiotic alone nor rBPI₂₁ alone had any effect on mortality other than to slightly retard the death rate. However, the concurrent administration of rBPI₂₁ with low-dose gentamicin

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significantly increased survival ($p < 0.5$ vs. vehicle). Concurrent administration of rBPI₁₁ with high-dose gentamicin dramatically increased survival ($p < 0.001$ vs. vehicle, and $p < 0.05$ vs. 0.03 mg/kg gentamicin with rBPI₁₁), protecting all but two of the mice from the lethal effects of bacterial challenge. The results clearly indicate synergism between gentamicin and BPI protein product. The greater synergistic effect of the concurrent administration of BPI protein product with gentamicin, compared to the effect of the concurrent administration of BPI protein product with cefamandole, may be related to the fact that aminoglycosides, which inhibit protein synthesis, have a different mechanism of action than BPI protein product.

Example 10

EFFECT OF BPI PROTEIN PRODUCT *IN VITRO* ON ANTIBIOTIC SUSCEPTIBILITY OF CEFTRIAXONE-RESISTANT GRAM-NEGATIVE ORGANISMS

The ability of BPI protein product, BPI₁₁, to reverse the resistance of a variety of gram-negative organisms to ceftriaxone (Roche Laboratories) was evaluated *in vitro*.

The strain of gram-negative bacteria to be tested was grown overnight at 37°C on Trypticase soy agar (TSA) plates. Colonies from the plate were then inoculated into nutrient broth (or triethanolamine-buffered minimal salts medium), grown overnight to stationary growth phase, diluted 1:10 in fresh medium, and grown to mid-late logarithmic growth phase (3 to 4 hours at 37°C) to an approximate concentration of 6 to 10×10^8 organisms/mL. Organism counts were performed by making serial dilutions, plating in triplicate on TSA plates, incubating at 37°C overnight, and counting colonies by visual inspection. Following development of a standard curve, counts were made by measuring OD₅₄₀ and confirming by plating. The bacteria were sedimented by centrifugation at 6,000g for 10 min. and resuspended in sterile saline to the desired concentration.

Ceftriaxone solutions were prepared from standard powder (Roche Laboratories). Solutions of BPI protein product were prepared from rBPI₁₁, Hanks solution, vitamin-free caseamino acid, and TRIS-HCl buffer, pH 7.0. Broth MIC

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studies were performed as follows: U-bottom, disposable, microtiter array plates (Dynatech) containing 100 μ L/well of broth and serial dilutions of 1:1 proportion rBPI₂₁ and ceftriaxone were prepared using a multichannel pipetting instrument, inoculated with organisms diluted to approximately 2.5×10^8 CFU/ml, and
5 incubated overnight at 37°C. Media controls containing no organism and growth controls containing organisms but no ceftriaxone or rBPI₂₁ were also prepared. The MIC was determined as the lowest drug concentration (μ g/mL) that inhibited bacterial growth. Agar MIC studies were performed as follows: The organism was grown in Mueller-Hinton broth (Difco Laboratories) overnight at 37°C,
10 transferred and grown to logarithmic phase, counted by optical densitometry, diluted with buffer and rBPI₂₁ and ceftriaxone, and incubated for 30 minutes at 37°C. After the 30 minutes of incubation, samples were serially diluted in sterile saline and plated onto TSA agar for bacterial counts after overnight incubation at 37°C.

15 A variety of different bacterial species were tested, including *Pseudomonas*, *Enterobacter*, *Citrobacter*, *Klebsiella*, and *Escherichia* species. A summary of the results is displayed in Table 11 below, which reports the minimum inhibitory concentration (MIC) of BPI protein product alone, the MIC of ceftriaxone alone, and the MIC of both agents together (in a fixed 1:1 proportion
20 of rBPI₂₁ and ceftriaxone). The experiments were replicated, and each number given in the table represents the highest, or worst-case, MIC for the group.

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TABLE 11
Synergistic Effect of rBPI_n and Ceftriaxone
on Ceftriaxone-Resistant Organisms

Organism	MIC of BPI alone ($\mu\text{g/mL}$)	MIC of Ceftriaxone alone ($\mu\text{g/mL}$)	MIC of BPI + Ceftriaxone ($\mu\text{g/mL}$)
PA 589	64	> 64	4
PA 631	> 64	> 64	4
PA 672	32	32	2
PA 677	> 64	> 64	4
EA 658	> 64	> 64	4
CF 595	4	> 64	1
CF 596	4	> 64	1
CF 597	> 64	> 64	8
CF 598	8	32	2
CF 642	4	16	8
CF 661	32	16	1
KP 601	64	> 64	8
EC 004	> 64	> 64	< 0.1
EC 600	> 64	32	8
EC 664	64	> 64	32
ECL 03	16	> 64	16
ECL 05	16	64	16
ECL 07	> 64	4	16
ECL 13	> 64	> 64	16
ECL 14	16	> 64	32
ECL 15	> 64	> 64	32
ECL 19	> 64	> 64	32
PA 001	> 64	> 64	8
PA 003	> 64	> 64	8
PA 004	> 64	64	16
PA 005	> 64	> 64	64
PA 012	> 64	> 64	16
PA 014	> 64	> 64	8

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TABLE II Synergistic Effect of rBPI ₂₁ and Ceftriaxone on Ceftriaxone-Resistant Organisms			
Organism	MIC of BPI alone (μ g/mL)	MIC of Ceftriaxone alone (μ g/mL)	MIC of BPI + Ceftriaxone (μ g/mL)
PA 017	> 64	> 64	32
PA 023	> 64	> 64	32
PA 026	64	64	8
PA 027	64	64	8
PA 028	> 64	> 64	16

PA = *Pseudomonas aeruginosa*EA = *Enterobacter aerogenes*CF = *Citrobacter freundii*5 KP = *Klebsiella pneumoniae*EC = *Escherichia coli*ECL = *Enterobacter cloacae*

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Checkerboard synergy studies on selected strains were performed as follows. Microplates were prepared using the following organization: Column 1, control organisms (growth control); Columns 2 through 9, serial dilutions of ceftriaxone; Column 10, ceftriaxone alone; Column 11, rBPI₂₁ alone; Column 12, control media; Rows 1 through 8, serial dilutions of rBPI₂₁. Columns 2 through 9 thus contained a serial array of various proportions of ceftriaxone and BPI concentrations. All wells, except the media control, were inoculated with suspended organisms and incubated overnight at 37°C. Turbidity was recorded at 24 and 48 hours.

Results of a representative synergy study of the effects of the concurrent administration of rBPI₂₁ and ceftriaxone on ceftriaxone-resistant *E. coli* are shown in the checkerboard in Table 12 below. In this checkerboard assay, rBPI₂₁ and ceftriaxone were each serially diluted to concentrations ranging from 100 µg/ml to 0.8 µg/ml. These results show that the concurrent administration of both agents is synergistic. There was uniform growth in all growth control wells (containing bacteria, no rBPI₂₁ and no antibiotic), while there was no growth in the media control wells (containing no bacteria). There was uniform growth in all ceftriaxone control wells (bacteria with ceftriaxone alone). In the BPI control wells (bacteria with rBPI₂₁ alone) there was growth in the wells with a concentration of 6.2 µg/ml rBPI₂₁ or less, and no growth at higher concentrations of BPI protein product.

Survival (or kill) curve studies were performed as follows. Tubes containing (1) media alone, (2) media plus ceftriaxone, (3) media plus rBPI₂₁, and (4) media plus ceftriaxone and rBPI₂₁, were prepared, inoculated with the desired organism, and incubated as described above. At 0, 1, 2, 4, 8 and 24 hours, diluted aliquots from each tube were plated, incubated and counted as described above. Growth curves over 24 hours (plots of log₁₀ CFU/mL versus time) were constructed to demonstrate the dynamics of antibiotic-BPI protein product interactions on bacterial growth and survival.

Results of a representative killing curve study of the effects of the concurrent administration of rBPI₂₁ and ceftriaxone on ceftriaxone-resistant *E. coli* are shown in Figure 18. The filled square is the control (neither rBPI₂₁ nor

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5 ceftriaxone), the open square is rBPI₂₁ alone, the filled diamond is ceftriaxone alone, and the open diamond is the concurrent administration of ceftriaxone and rBPI₂₁. Ceftriaxone or rBPI₂₁ alone have some early bactericidal effect at 4-8 hours, but organism growth for both almost reaches that of the control curve by 24 hours (about a 1 log difference from control). In contrast, the concurrent administration of rBPI₂₁ and ceftriaxone produces a significantly greater bactericidal effect that is sustained at 24 hours (a more than 6 log difference from control).

TABLE 12
Checkerboard Synergy Study for Ceftriaxone-Resistant *E. coli* Using Ceftriaxone and rBPI₂

	0.8 µg/mL Ceftriaxone	1.5 µg/mL Ceftriaxone	3.1 µg/mL Ceftriaxone	6.2 µg/mL Ceftriaxone	12.5 µg/mL Ceftriaxone	25 µg/mL Ceftriaxone	50 µg/mL Ceftriaxone	100 µg/mL Ceftriaxone
0.8 µg/mL BPI	+	+	0	0	0	0	0	0
1.5 µg/mL BPI	+	0	0	0	0	0	0	0
3.1 µg/mL BPI	0	0	0	0	0	0	0	0
6.2 µg/mL BPI	0	0	0	0	0	0	0	0
12.5 µg/mL BPI	0	0	0	0	0	0	0	0
25 µg/mL BPI	0	0	0	0	0	0	0	0
50 µg/mL BPI	0	0	0	0	0	0	0	0
100 µg/mL BPI	0	0	0	0	0	0	0	0

KEY:
+ Growth
0 No Growth

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Example 11**EFFECTS OF BPI PROTEIN PRODUCT
AND ANTIBIOTICS *IN VITRO* ON
PSEUDOMONAS SPECIES**

5 The effects of a BPI protein product, rBPI₂₁, on the antibiotic
susceptibility of clinical isolates of *Pseudomonas aeruginosa* and other
Pseudomonas species (from Baxter Microscan® library, Sacramento, CA)
was evaluated using Microscan® panel plates (Baxter Diagnostics, Inc.,
Deerfield, IL) that allow simultaneous determination of minimum inhibitory
10 concentrations for a number of different antibiotics. Control assays
confirmed that the formulation buffer for rBPI₂₁ had no effect on the
antibiotic susceptibility of various organisms.

 The antimicrobial susceptibility tests performed on the
Microscan® panel plates are miniaturizations of the broth dilution
15 susceptibility test. Antimicrobial agents are serially diluted in Mueller-
Hinton broth (supplemented with calcium and magnesium, or with sodium
chloride for oxacillin, or with thymidine phosphorylase for trimethoprim,
sulfamethoxazole and trimethoprim/ sulfamethoxazole) to concentrations
bridging the range of clinical interest. One well on the 96-well Microscan®
20 plate is a growth control well that contains dehydrated broth only. The
remaining wells contain dehydrated broth and antibiotic (or broth and
biochemical reagent indicator), which is rehydrated to the desired
concentration by inoculation of a standardized suspension of test organism.
The chromogenic biochemical agent indicators are used to identify and
25 characterize the species of bacteria based on detection of pH changes and
substrate utilization. After incubation overnight, the minimum inhibitory
concentration (MIC) of an antibiotic for the test organism is determined by
observing the well with the lowest concentration of the antibiotic that shows
inhibition of growth. Gram-negative organisms were tested using Neg
30 Combo Type 16, MIC Plus Type 2, or Neg Breakpoint Combo Type 9
panel plates (Microscan®, Baxter Diagnostics, Inc., Deerfield, IL). The

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5 concentrations of antibiotics tested in these panel plates are shown below in Tables 13, 14 and 15, respectively. The antibiotic susceptibility standards (interpretation of an MIC as resistant, intermediate or susceptible according to Microscan®'s NCCLS-derived standards) applicable to the gram-negative organisms tested in each panel plate appear in Tables 13A, 14A and 15A, respectively.

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TABLE 13 ANTIBIOTIC CONCENTRATIONS TESTED IN NEG COMBO TYPE 16 PANEL PLATE	
Antibiotic	Two-Fold Serial Dilutions Tested $\mu\text{g/ml}$
Amikacin	2-16
Ampicillin	2-16
Ampicillin/Sulbactam	8/4-16/8
Aztreonam	8-16
Cefazolin	2-16
Cefotaxime	4-32
Cefoxitin	2-16
Ceftazidime	2-16
Ceftriaxone	4-32
Cefuroxime	2-16
Ciprofloxacin	1-2
Gentamicin	1-4, 6
Imipenem	4-8
Ofloxacin	2-4
Piperacillin	8-64
Ticarcillin	8-64
Tobramycin	1-4, 6
Trimethoprim/Sulfamethoxazole	0.5/9.5, 2/38

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TABLE 13A
MICROSCAN NEG COMBO PANEL 16
ANTIBIOTIC SUSCEPTIBILITY RANGES
FOR GRAM-NEGATIVE BACTERIA

Antibiotic	MIC ($\mu\text{g/ml}$)		
	Resistant	Intermediate	Susceptible
Amikacin	> 16		≤ 16
Ampicillin ^s	> 16	16	≤ 8
Ampicillin/Sulbactam ^s	> 16/8	16/8	$\leq 8/4$
Aztreonam	> 16	16	≤ 8
Cefazolin	> 16	16	≤ 8
Cefotaxime	> 32	16-32	≤ 8
Cefoxitin	> 16	16	≤ 8
Ceftazidime	> 16	16	≤ 8
Ceftriaxone	> 32	16-32	≤ 8
Cefuroxime	> 16	16	≤ 8
Ciprofloxacin	> 2	2	≤ 1
Gentamicin	> 6	6	≤ 4
Imipenem	> 8	8	≤ 4
Ofloxacin	> 4	4	≤ 2
Piperacillin ^s	> 64	32-64	≤ 16
Piperacillin ^p	> 64		≤ 64
Ticarcillin ^s	> 64	32-64	≤ 16
Ticarcillin ^p	> 64		≤ 64
Tobramycin	> 6	6	≤ 4
Trimethoprim/Sulfamethoxazole	> 2/38		$\leq 2/38$

^s *Enterobacteriaceae* only^p *Pseudomonas* only

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TABLE 14
ANTIBIOTIC CONCENTRATIONS TESTED IN
MIC PLUS TYPE 2 PANEL PLATE

Antibiotic	Two-Fold Serial Dilutions Tested ($\mu\text{g/ml}$)
Amoxicillin/K Clavulanate	1/0.5-32/16
Ampicillin/Sulbactam	1/0.5-32/16
Azlocillin	64
Aztreonam	1-32
Carbenicillin	16-128
Cefamandole	4-32
Cefonicid	2-16
Cefoperazone	4-32
Cefotaxime	2-64
Cefotetan	4-32
Ceftazidime	1-32
Ceftizoxime	2-32
Ceftriaxone	2-64
Chloramphenicol	2-16
Ciprofloxacin	0.25-4
Imipenem	0.5-16
Mezlocillin	16-128
Netilmicin	2-16
Ticarcillin	16-128
Ticarcillin/K Clavulanate	16-128

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TABLE 14A
MICROSCAN MIC PLUS TYPE 2 ANTIBIOTIC
SUSCEPTIBILITY RANGES FOR GRAM-NEGATIVE BACTERIA

Antibiotic	MIC (µg/ml)		
	Resistant	Intermediate	Susceptible
Amoxicillin/K Clavulanate	≥ 32/16	16/8	≤ 8/4
Ampicillin/Sulbactam	≥ 32/16	16/8	≤ 8/4
Azlocillin ^a	> 64		≤ 64
Aztreonam	≥ 32	16	≤ 8
Carbenicillin ^a	≥ 64	32	≤ 16
Carbenicillin ^b	> 128		≤ 128
Cefamandole	≥ 32	16	≤ 8
Cefonicid	> 16	16	≤ 8
Cefoperazone	> 32	32	≤ 16
Cefotaxime	≥ 64	16-32	≤ 8
Cefotetan	> 32	32	≤ 16
Ceftazidime	≥ 32	16	≤ 8
Ceftizoxime	> 32	16-32	≤ 8
Ceftriaxone	≥ 64	16-32	≤ 8
Chloramphenicol	> 16	16	≤ 8
Ciprofloxacin	≥ 4	2	≤ 1
Imipenem	≥ 16	8	≤ 4
Mezlocillin ^a	≥ 128	32-64	≤ 16
Mezlocillin ^b	≥ 128		≤ 64
Netilmicin	> 16	16	≤ 8
Ticarcillin ^a	≥ 128	32-64	≤ 16
Ticarcillin ^b	≥ 128		≤ 64
Ticarcillin/K Clavulanate ^a	≥ 128	32-64	≤ 16
Ticarcillin/K Clavulanate ^b	≥ 128		≤ 64

^a Enterobacteriaceae only

^b Pseudomonas only

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TABLE 15
ANTIBIOTIC CONCENTRATIONS TESTED IN
NEG BREAKPOINT COMBO TYPE 9 PANEL PLATE

Antibiotic	Dilutions Tested (µg/ml)
Nitrofurantoin	32 & 64
Cephalothin	8 & 16
Ampicillin	8 & 16
Ofloxacin	2 & 4
Ticarcillin	16 & 64
Piperacillin	16 & 64
Mezlocillin	16 & 64
Tetracycline	4 & 8
Ampicillin/Sulbactam	8/4 & 16/8
Amoxicillin/K Clavulanate	8/4 & 16/8
Ticarcillin/K Clavulanate	16 & 64
Gentamicin	4 & 8
Tobramycin	4 & 8
Amikacin	16 & 32
Ciprofloxacin	1 & 2
Imipenem	4 & 8
Cefazolin	8 & 16
Cefamandole	8 & 16
Cefuroxime	8 & 16
Cefotetan	16 & 32
Cefoxitin	8 & 16
Aztreonam	8 & 16
Ceftriaxone	8 & 32
Cefazidime	8 & 16
Cefoperazone	16 & 32
Cefotaxime	8 & 32
Chloramphenicol	8 & 16
Trimethoprim/Sulfamethoxazole	2/38 & 8/152
Norfloxacin	4 & 8
Cinoxacin	16
Trimethoprim	8
Sulfamethoxazole	256

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TABLE 15A
MICROSCAN NEG BREAKPOINT COMBO TYPE 9
ANTIBIOTIC SUSCEPTIBILITY RANGES
FOR GRAM-NEGATIVE BACTERIA

Antibiotic	MIC (µg/ml)		
	Resistant	Intermediate	Susceptible
Nitrofurantoin	> 64	64	≤ 32
Cephalothin	> 16	16	≤ 8
Ampicillin [†]	> 16	16	≤ 8
Ofloxacin	> 4	4	≤ 2
Ticarcillin [†]	> 64	64	≤ 16
Ticarcillin [†]	> 64		≤ 64
Piperacillin [†]	> 64	64	≤ 16
Piperacillin [†]	> 64		≤ 64
Mezlocillin [†]	> 64	64	16
Mezlocillin [†]	> 64		≤ 64
Tetracycline	> 8	8	≤ 4
Ampicillin/Sulbactam [†]	> 16/8	16/8	≤ 8/4
Amoxicillin/K Clavulanate	> 16/8	16/8	≤ 8/4
Ticarcillin/K Clavulanate [†]	> 64	64	≤ 16
Ticarcillin/K Clavulanate [†]	> 64		≤ 64
Gentamicin	> 8	8	≤ 4
Tobramycin	> 8	8	≤ 4
Amikacin	> 32	32	≤ 16
Ciprofloxacin	> 2	2	≤ 1
Imipenem	> 8	8	≤ 4
Cefazolin	> 16	16	≤ 8
Cefamandole	> 16	16	≤ 8
Cefuroxime	> 16	16	≤ 8
Cefotetan	> 32	32	≤ 16
Cefoxitin	> 16	16	≤ 8
Aztreonam	> 16	16	≤ 8
Ceftriaxone	> 32	32	≤ 8
Ceftazidime	> 16	16	≤ 8
Cefoperazone	> 32	32	≤ 16
Cefotaxime	> 32	32	≤ 8
Chloramphenicol	> 16	16	≤ 8
Trimethoprim/Sulfamethoxazole	> 8/152	8/152	≤ 2/38
Norfloxacin	> 8	8	≤ 4
Cinoxacin	> 16		≤ 16
Trimethoprim	> 8		≤ 8
Sulfamethoxazole	> 256		≤ 256

[†] Enterobacteriaceae only
Pseudomonas only

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For each experimental run, the following procedure was performed: The organism was streaked onto 5% sheep blood agar plates (Remel, Lenexa, Kansas) and incubated for 18-24 hours overnight. Well-isolated colonies from the plates were emulsified in 3 ml of sterile Inoculum Water (catalog no. B1015-2, MicroScan® system, Baxter Diagnostics, Inc., Deerfield, IL) to a final turbidity equivalent to 0.5 McFarland Barium Sulfate standard. This cell suspension was vortexed for 2 to 3 seconds and 100 μ l was transferred to glass tubes containing 25 ml of Inoculum Water with Pluronic-D (catalog no. B1015-7, MicroScan® system, Baxter Diagnostics, Inc., Deerfield, IL) (hereinafter "Pluronic Inoculum Water"), or 25 ml of Pluronic Inoculum Water into which rBPI₁₁ (in formulation buffer) had been diluted to the desired concentration between 0.5 to 64 μ g/ml rBPI₁₁.

The 25 ml of this inoculum containing rBPI₁₁ was mixed by inversion and poured into a tray. The inoculum was drawn up into a manual 96-well pipetting system (RENOK® rehydrator-inoculator system, Baxter Health Care Corporation, West Sacramento, CA) designed for use with the Microscan® panel plates, and 110 μ l of the inoculum was delivered to each well of a Microscan® Neg Combo Type 16 panel plate. When added to the wells, this inoculum achieves a final bacterial concentration of 4×10^4 to 7×10^5 CFU/ml. The panel plates were then incubated at 35°C for 15-24 hours and read visually for cell growth.

No growth was defined as a slight whiteness in the well or a clear broth. Growth appeared as turbidity which could take the form of a white haze throughout the well, a white button in the center of the well, or a fine granule growth throughout the well. All wells were read against a black indirectly lighted background. Visual results of the biochemical reactions were read into a database for bacterial identification. The MICs for each antibiotic tested were determined by identifying the lowest concentration of antibiotic which inhibited visible growth.

The clinical isolates of *Pseudomonas aeruginosa* and other *Pseudomonas* species were tested using the Neg Combo Type 16 panel plate. Tables 16, 17 and 18 below display a summary of the results of the antibiotic

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screening panels, reported for each strain tested as the MIC of the tested antibiotics at the various concentrations of rBPI₂₁ indicated. Results are reported for each strain tested, but susceptibility data is listed for only those antibiotics for which BPI protein product altered susceptibility. The antibiotic susceptibility standards (interpretation of an MIC as resistant, intermediate or susceptible according to Microscan®'s NCCLS-derived standards) applicable to the organism tested appear in Table 13A. Stars after the antibiotic name in the "antibiotic tested" column indicate whether rBPI₂₁ reversed the resistance of that organism to the antibiotic tested (two stars) or converted an indifferent MIC into a susceptible MIC (one star). These data show that BPI protein product reversed resistance to ticarcillin, cefazolin, ceftazidime, cefuroxime, ofloxacin, aztreonam, piperacillin, and amikacin for some strains of *P. aeruginosa* and increased the susceptibility of some *P. aeruginosa* strains to ticarcillin, aztreonam, imipenem, piperacillin, ofloxacin, ceftazidime, amikacin, ceftriaxone, cefotaxime, cefuroxime, tobramycin, ciprofloxacin, trimethoprim/sulfamethoxazole, gentamicin, and cefazolin. BPI protein product reversed resistance of some *P. cepacia* strains to cefazolin, ceftazidime, cefuroxime, ceftriaxone, ticarcillin, and increased the susceptibility of some *P. cepacia* strains to ampicillin, ticarcillin, piperacillin, cefazolin, ceftazidime, cefuroxime, ceftriaxone, ampicillin/sulbactam, cefotaxime, gentamicin, tobramycin, and amikacin. BPI protein product reversed resistance of *Xanthomonas maltophilia* to trimethoprim/sulfamethoxazole, piperacillin and amikacin, and increased susceptibility to ciprofloxacin.

Tables 16, 17 and 18 also show the presence or absence of bacterial growth in the growth control wells, which contained varying concentrations of rBPI₂₁ alone without antibiotic. "G" indicates growth, while "NG" indicates no growth. These results indicate that rBPI₂₁ at a concentration of 32 µg/ml has direct bactericidal/growth inhibitory effects on some of the tested *Pseudomonas* isolates.

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TABLE 16
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Pseudomonas aeruginosa*

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 8 μg/mL rBPI ₂₁	With 32 μg/mL rBPI ₂₁
19610	BPI	G	G	G
	Ticarcillin**	> 64	64	64
	Aztreonam*	16	< 8	< 8
	Piperacillin	32	16	< 8
	Ofloxacin**	> 4	4	< 2
	Ceftazidime	8	8	4
	Amikacin	16	8	8
18433	BPI	G	G	NG
	Aztreonam*	16	< 8	--
	Ceftriaxone	> 32	32	--
	Cefotaxime	> 32	32	--
	Cefuroxime*	16	4	--
	Tobramycin	2	< 1	--
	Amikacin	16	8	--
12892	BPI	G	G	G
	Ciprofloxacin	> 2	> 2	2
	Ofloxacin	> 4	> 4	4
	Ceftazidime	4	4	< 2
	Cefotaxime	> 32	> 32	32
	Amikacin	16	16	8

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TABLE 16
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Pseudomonas aeruginosa*

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 8 μg/mL rBPI ₂₁	With 32 μg/mL rBPI ₂₁
19054	BPI	G	G	G
	Trimethoprim/Sulfamethoxazole	>2	>2	2
	Ticarcillin	64	16	16
	Ceftriaxone*	16	32	8
	Gentamicin	4	4	2
	Amikacin	16	8	4
	Cefazolin	>16	>16	16
19672	BPI	G	G	G
	Tobramycin	>6	>6	6
19551	BPI	G	G	G
	Aztreonam**	>16	<8	<8
	Piperacillin**	>64	<8	<8
	Ceftriaxone	32	16	16
	Gentamicin	4	<1	<1
	Tobramycin	2	<1	<1
	Amikacin**	>16	<2	<2

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TABLE 16
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Pseudomonas aeruginosa*

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ²¹	With 8 μg/mL rBPI ²¹	With 32 μg/mL rBPI ₂₁
19660	BPI	G	G	NG
	Trimethoprim/Sulfamethoxazole	>2	2	—
	Ceftriaxone*	16	8	—
	Cefotaxime	16	16	—
Opal ^b	BPI	G	G	G
	Ampicillin*	>16	>16	4
	Trimethoprim/Sulfamethoxazole	>2	>2	2
	Ticarcillin	16	16	<8
	Cefazolin**	>16	>16	<2
	Cefoxitin**	>16	>16	<2
	Cefuroxime**	>16	>16	<2
	Ceftriaxone*	16	8	<4
	Ampicillin/Sulbactam*	>16	>16	<8
	Cefotaxime*	16	8	<4

* The Microscan® worksheet did not supply antibiotic susceptibility ranges, but there was reversal of antibiotic indifference according to NCCLS standards, Publication M7-A3, Table 2 (1993).

^b Strain 12.4.4, provided by S.M. Opal, Brown University, Providence, RI.

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TABLE 17
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Pseudomonas aeruginosa*

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
N113-100	BPI	G	G	G
	Ofloxacin*	4	4	<2
	Gentamicin	>6	6	6
	Tobramycin	2	2	<1
N113-101	BPI	G	G	G
	Aztreonam	>16	>16	<8
	Ofloxacin*	4	4	<2
	Gentamicin	>6	6	6
	Tobramycin	4	2	<1
N113-102	BPI	G	G	G
	Piperacillin	64	<8	<8
	Ceftriaxone*	32	16	8
	Cefotaxime	32	16	16
	Gentamicin	4	4	<1
	Amikacin	8	8	<2
N113-103	BPI	G	G	G
	Ticarcillin	16	16	32
	Amikacin	4	8	4

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TABLE 17 EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON <i>Pseudomonas aeruginosa</i>				
Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
N113-104	BPI	G	G	G
	Ceftazidime	<2	<2	4
	Ceftriaxone*	16	8	8
N113-105	BPI	G	G	G
	Ticarcillin	64	32	32
	Aztreonam*	16	<8	<8
	Ceftriaxone	32	32	16
	Cefotaxime	32	32	16
	Gentamicin	2	2	<1
N113-106	BPI	G	G	G
	Ticarcillin	16	16	<8
	Ceftriaxone*	32	16	8
	Cefotaxime*	16	16	8
	Gentamicin	4	2	2
	Amikacin	8	4	4
N113-107	BPI	G	G	G
	Ticarcillin	> 64	> 64	64
	Ciprofloxacin*	2	2	<1
	Ceftazidime	8	4	4
	Ceftriaxone	32	32	> 32
	Imipenem	> 8	8	> 8

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TABLE 18
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Pseudomonas* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 8 μg/mL rBPI ₂₁	With 32 μg/mL rBPI ₂₁
31142 (<i>P. cepacia</i>)	BPI	G	G	NG
	Ampicillin	> 16	16	--
	Ticarcillin	32	< 8	--
	Piperacillin	32	< 8	--
	Cefazolin**	> 16	8	--
	Cefoxitin**	> 16	< 2	--
	Cefuroxime**	> 16	< 2	--
	Ceftriaxone**	> 32	8	--
	Ampicillin/ Sulbactam*	16	< 8	--
	Cefotaxime*	32	< 4	--
	Gentamicin*	6	< 1	--
	Tobramycin	2	< 1	--
	Amikacin	8	< 2	--
12122 (<i>P. cepacia</i>)	BPI	G	G	G
	Ticarcillin**	> 64	64	32
	Piperacillin	16	< 8	< 8
	Cefuroxime	4	< 2	< 2
	Amikacin	4	4	< 2

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TABLE 18
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Pseudomonas* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (µg/mL)		
		With 0 µg/mL rBPI ₂₁	With 8 µg/mL rBPI ₂₁	With 32 µg/mL rBPI ₂₁
17211 (<i>Xanthamonas</i> <i>malisophila</i>)	BPI	G	G	NG
	Trimethoprim/ Sulfamethoxazole**	> 2	2	--
	Piperacillin**	> 64	16	--
	Amikacin**	> 16	16	--
	Ciprofloxacin	2	2	--

* The Microscan® worksheet did not supply antibiotic susceptibility ranges for this organism, but there was reversal of antibiotic indifference according to NCCLS standards, Publication M7-A3, Table 2 (1993).

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Example 12**EFFECTS OF BPI PROTEIN PRODUCT
AND ANTIBIOTICS *IN VITRO* ON STRAINS OF *E. COLI***

5 The effect of a BPI protein product, rBPI₂₁, on the antibiotic
susceptibility of various strains of *E. coli* was evaluated in the Microscan®
antibiotic susceptibility screening assay of Example 11 using the Neg
Combo Type 16 panel plate. The direct growth inhibitory effect of rBPI₂₁
on these strains was also evaluated in the same assay. Assays were
conducted on clinical isolates of *E. coli* (from Baxter Microscan® library,
10 Sacramento, CA).

A summary of the results of the antibiotic screening panels,
reported as MICs (µg/ml) of the antibiotic tested, is shown in Table 19
below. Results are reported for each strain tested, but susceptibility data is
listed for only those antibiotics for which BPI protein product altered
15 susceptibility. The antibiotic susceptibility standards (interpretation of an
MIC as resistant, intermediate or susceptible according to Microscan®'s
NCCLS-derived standards) applicable to the organism tested appear in
Table 13A. These results show that BPI protein product reversed the
resistance of some strains to cefazolin and increased the susceptibility of
20 other strains to ampicillin, cefuroxime, cefazolin, amikacin, and ceftiofur.

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TABLE 19
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Escherichia coli*

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (µg/mL)		
		With 0 µg/mL rBPI ₂₁	With 4 µg/mL rBPI ₂₁	With 16 µg/mL rBPI ₂₁
19536	Ampicillin*	16	16	8
	Cefuroxime	4	8	4
F101-309	Cefoxitin	4	<2	4
	Cefuroxime	8	4	4
19612	Cefazolin**	> 16	> 16	8
	Amikacin	16	8	8
	Cefoxitin	<2	4	<2
17164	Ampicillin	8	8	4
	Cefoxitin	8	8	4
19522	Cefazolin	8	8	4
	Cefoxitin	> 16	> 16	16

* The Microscan® worksheet did not supply antibiotic susceptibility ranges for this organism, but there was reversal of antibiotic indifference according to NCCLS standards, Publication M7-A3, Table 2 (1993).

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Example 13**EFFECTS OF BPI PROTEIN PRODUCT
AND ANTIBIOTICS *IN VITRO* ON *CITROBACTER* SPECIES**

5 The effect of a BPI protein product, rBPI₂₁, on the antibiotic susceptibility of various *Citrobacter* species was evaluated in the Microscan® antibiotic susceptibility screening assay of Example 11 using the Neg Combo Type 16 panel plate. The direct growth inhibitory effect of rBPI₂₁ on these strains was also evaluated in the same assay. Assays were conducted on clinical isolates of *Citrobacter* species (from Baxter
10 Microscan® library, Sacramento, CA).

A summary of the results of the antibiotic screening panels, reported as MICs (µg/ml) of the antibiotic tested, is shown in Table 20 below. Results are reported for each strain tested, but susceptibility data is listed for only those antibiotics for which BPI protein product altered
15 susceptibility. The antibiotic susceptibility standards (interpretation of an MIC as resistant, intermediate or susceptible according to Microscan®'s NCCLS-derived standards) applicable to the organism tested appear in Table 13A. These results show that BPI protein product increased the susceptibility of the tested *Citrobacter* species to aztreonam, cefotaxime,
20 tobramycin, amikacin, cefuroxime, ampicillin, ticarcillin, piperacillin, and cefuroxime.

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TABLE 20 EFFECTS OF rBPI ₂₁ ± ANTIBIOTICS ON <i>Citrobacter</i> SPECIES				
Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (µg/mL)		
		With 0 µg/mL rBPI ₂₁	With 4 µg/mL rBPI ₂₁	With 16 µg/mL rBPI ₂₁
18419 (<i>C. freundii</i>)	Azureonam	> 16	> 16	16
	Cefotaxime	> 32	32	32
	Tobramycin	4	2	< 1
	Amikacin	4	< 2	4
18420 (<i>C. freundii</i>)	Cefuroxime	> 16	> 16	16
	Amikacin	> 2	> 2	> 2
F052-007 (<i>C. diversus</i>)	Ampicillin	> 16	> 16	16
	Ticarcillin	> 64	> 64	64
	Piperacillin*	32	16	< 8
	Cefuroxime	8	4	4

* The Microscan® worksheet did not supply antibiotic susceptibility ranges for this organism, but there was reversal of antibiotic indifference according to NCCLS standards, Publication M7-A3, Table 2 (1993).

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Example 14**EFFECTS OF BPI PROTEIN PRODUCT
AND ANTIBIOTICS *IN VITRO* ON *KLEBSIELLA* SPECIES**

The effect of a BPI protein product, rBPI₂₁, on the antibiotic susceptibility of various *Klebsiella* species was evaluated in the Microscan® antibiotic susceptibility screening assay of Example 11 using the Neg Combo Type 16 panel plate. The direct growth inhibitory effect of rBPI₂₁ on these strains was also evaluated in the same assay. Assays were conducted on clinical isolates of *Klebsiella* species (from Baxter Microscan® library, Sacramento, CA).

A summary of the results of the antibiotic screening panels, reported as MICs (µg/ml) of the antibiotic tested, is shown in Table 21 below. Results are reported for each strain tested, but susceptibility data is listed for only those antibiotics for which BPI protein product altered susceptibility. The antibiotic susceptibility standards (interpretation of an MIC as resistant, intermediate or susceptible according to Microscan®'s NCCLS-derived standards) applicable to the organism tested appear in Table 13A. These results show that BPI protein product reversed the resistance of one strain of *K. pneumoniae* to trimethoprim/sulfamethoxazole and increased the susceptibility of the tested species to cefoxitin, ampicillin/sulbactam, trimethoprim/sulfamethoxazole, cefazolin, and cefuroxime.

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TABLE 21
EFFECTS OF rBPI₁₁ ± ANTIBIOTICS ON *Klebsiella* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (µg/mL)		
		With 0 µg/mL rBPI ₁₁	With 4 µg/mL rBPI ₁₁	With 16 µg/mL rBPI ₁₁
19645 (<i>K. pneumoniae</i>)	Cefoxitin	8	<2	4
	Ampicillin/ Sulbactam	>16	>16	>16
18427 (<i>K. pneumoniae</i>)	Cefazolin	<2	4	<2
	Ampicillin/ Sulbactam*	16	16	<8
16135 (<i>K. pneumoniae</i>)	Trimethoprim/ Sulfamethoxazole**	>2	>2	<0.5
	Cefazolin*	16	16	8
30434 (<i>K. oxytoca</i>)	Cefazolin*	16	16	8
	Cefoxitin	4	<2	4
	Cefuroxime	8	4	8
	Ampicillin/ Sulbactam	<8	16	<8

* The Microscan® worksheet did not supply antibiotic susceptibility ranges for this organism, but there was reversal of antibiotic indifference according to NCCLS standards, Publication M7-A3, Table 2 (1993).

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Example 15**EFFECTS OF BPI PROTEIN PRODUCT
AND ANTIBIOTICS IN VITRO ON ENTEROBACTER SPECIES**

The effect of a BPI protein product, rBPI₂₁₁, on the antibiotic susceptibility of various *Enterobacter* species was evaluated in the Microscan® antibiotic susceptibility screening assay of Example 11 using the Neg Combo Type 16 panel plate. The direct growth inhibitory effect of rBPI₂₁₁ on these strains was also evaluated in the same assay. Assays were conducted on clinical isolates of *Enterobacter* species (from Baxter Microscan® library, Sacramento, CA).

A summary of the results of the antibiotic screening panels, reported as MICs ($\mu\text{g/ml}$) of the antibiotic tested, is shown in Table 22 below. Results are reported for each strain tested, but susceptibility data is listed for only those antibiotics for which BPI protein product altered susceptibility. The antibiotic susceptibility standards (interpretation of an MIC as resistant, intermediate or susceptible according to Microscan®'s NCCLS-derived standards) applicable to the organism tested appear in Table 13A. These results show that BPI protein product reversed the resistance of one strain of *E. cloacae* to ticarcillin, cefuroxime, ceftazidime and cefotaxime. BPI protein product also increased the susceptibility of some *Enterobacter* species to ticarcillin, aztreonam, piperacillin, ciprofloxacin, cefotaxime, trimethoprim/sulfamethoxazole, cefuroxime, ceftazidime, ceftriaxone, and ampicillin/sulbactam.

TABLE 22
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Enterobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
19565 (<i>E. aerogenes</i>)	Ticarcillin	64	> 64	64
	Aztreonam*	16	< 8	< 8
	Piperacillin	> 64	> 64	64
	Ciprofloxacin*	2	< 1	< 1
	Cefotaxime*	16	8	8
19626 (<i>E. aerogenes</i>)	Piperacillin	> 64	64	64
	Ticarcillin	> 64	> 64	> 64
19625 (<i>E. aerogenes</i>)	Trimethoprim/Sulfamethoxazole*	2	< 0.5	< 0.5
	Piperacillin	> 64	> 64	> 64
19680 (<i>E. cloacae</i>)	Ticarcillin**	> 64	> 64	< 8
	Aztreonam	> 16	16	< 8
	Piperacillin*	64	32	< 8
	Cefuroxime**	> 16	> 16	8
	Ceftazidime**	> 32	32	< 4
	Ceftriaxone	> 16	> 16	16
	Ampicillin/Sulbactam	> 16	> 16	16
	Cefotaxime**	> 32	> 32	< 4
19686 (<i>E. cloacae</i>)	Ceftriaxone	32	32	16
	Piperacillin	> 64	> 64	> 64

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Example 16**EFFECTS OF BPI PROTEIN PRODUCT
AND ANTIBIOTICS *IN VITRO* ON *SERRATIA MARCESCENS***

5 The effect of a BPI protein product, rBPI₂₁, on the antibiotic
susceptibility of *Serratia marcescens* was evaluated in the Microscan®
antibiotic susceptibility screening assay of Example 11 using the Neg
Combo Type 16 panel plate. The direct growth inhibitory effect of rBPI₂₁
on these strains was also evaluated in the same assay. Assays were
conducted on clinical isolates of *Serratia marcescens* (from Baxter
10 Microscan® library, Sacramento, CA).

A summary of the results of the antibiotic screening panels,
reported as MICs (µg/ml) of the antibiotic tested, is shown in Table 23
below. Results are reported for each strain tested, but susceptibility data is
listed for only those antibiotics for which BPI protein product altered
15 susceptibility. The antibiotic susceptibility standards (interpretation of an
MIC as resistant, intermediate or susceptible according to Microscan®'s
NCCLS-derived standards) applicable to the organism tested appear in
Table 13A. These results show that BPI protein product reversed the
resistance of some strains to ceftazidime and cefotaxime, and increased the
20 susceptibility of other strains to piperacillin, cefoxitin, ceftazidime,
ceftriaxone, cefotaxime, tobramycin, ampicillin/sulbactam, and ampicillin.

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TABLE 23 EFFECTS OF rBPI ₂₁ ± ANTIBIOTICS ON <i>Serratia marcescens</i>				
Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (µg/mL)		
		With 0 µg/mL rBPI ₂₁	With 8 µg/mL rBPI ₂₁	With 32 µg/mL rBPI ₂₁
19646	Piperacillin	> 64	> 64	32
	Cefoxitin	16	> 16	8
	Ceftazidime**	> 16	> 16	8
	Ceftriaxone*	32	16	8
	Cefotaxime**	> 32	32	< 4
	Tobramycin*	6	6	4
19647	Piperacillin*	> 64	64	16
	Cefoxitin	8	8	4
	Ceftazidime*	16	16	4
	Ceftriaxone	8	< 4	< 4
	Ampicillin/ Sulbactam	> 16	> 16	16
	Cefotaxime	8	< 4	< 4
	Tobramycin	2	2	< 1
18443	Ampicillin*	16	16	8
	Ampicillin/ Sulbactam	> 16	> 16	16

* The Microscan® worksheet did not supply antibiotic susceptibility ranges for this organism, but there was reversal of antibiotic indifference according to NCCLS standards, Publication M7-A3, Table 2 (1993).

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Example 17**EFFECTS OF BPI PROTEIN PRODUCT
AND ANTIBIOTICS *IN VITRO* ON *PROTEUS MIRABILIS***

5 The effect of a BPI protein product, rBPI₂₁, on the antibiotic
susceptibility of *Proteus mirabilis* was evaluated in the Microscan®
antibiotic susceptibility screening assay of Example 11 using the Neg
Combo Type 16 panel plate. The direct growth inhibitory effect of rBPI₂₁
on these strains was also evaluated in the same assay. Assays were
10 conducted on clinical isolates of *Proteus mirabilis* (from Baxter Microscan®
library, Sacramento, CA).

A summary of the results of the antibiotic screening panels,
reported as MICs (µg/ml) of the antibiotic tested, is shown in Table 24
below. Results are reported for each strain tested, but susceptibility data is
15 listed for only those antibiotics for which BPI protein product altered
susceptibility. The antibiotic susceptibility standards (interpretation of an
MIC as resistant, intermediate or susceptible according to Microscan®'s
NCCLS-derived standards) applicable to the organism tested appear in
Table 13A. These results show that BPI protein product increased the
20 susceptibility of some strains to trimethoprim/sulfamethoxazole, cefazolin,
cefoxitin, imipenem, tobramycin, and amikacin.

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TABLE 24 EFFECTS OF rBPI ₂₁ ± ANTIBIOTICS ON <i>Proteus mirabilis</i>				
Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (µg/mL)		
		With 0 µg/mL rBPI ₂₁	With 8 µg/mL rBPI ₂₁	With 32 µg/mL rBPI ₂₁
19593	Trimethoprim/ Sulfamethoxazole*	2	<0.5	<0.5
	Cefazolin	4	<2	4
	Cefoxitin	8	<2	<2
	Imipenem*	8	<4	<4
	Tobramycin	2	2	<1
	Amikacin	8	4	4
F231-129	Cefoxitin	4	<2	4
	Imipenem*	8	<4	<4
	Amikacin	4	<2	<2

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Example 18**EFFECTS OF BPI PROTEIN PRODUCT
AND ANTIBIOTICS *IN VITRO* ON *PROVIDENCIA* SPECIES**

5 The effect of a BPI protein product, rBPI₂₁, on the antibiotic
susceptibility of various *Providencia* species was evaluated in the
Microscan® antibiotic susceptibility screening assay of Example 11 using
the Neg Combo Type 16 panel plate. The direct growth inhibitory effect
of rBPI₂₁ on these strains was also evaluated in the same assay. Assays
10 were conducted on clinical isolates of *Providencia* species (from Baxter
Microscan® library, Sacramento, CA).

A summary of the results of the antibiotic screening panels,
reported as MICs (µg/ml) of the antibiotic tested, is shown in Table 25
below. Results are reported for each strain tested, but susceptibility data is
15 listed for only those antibiotics for which BPI protein product altered
susceptibility. The antibiotic susceptibility standards (interpretation of an
MIC as resistant, intermediate or susceptible according to Microscan®'s
NCCLS-derived standards) applicable to the organism tested appear in
Table 13A. These results show that BPI protein product reversed the
20 resistance of *P. stuartii* to cefazolin and cefuroxime, and increased its
susceptibility to piperacillin, ceftazidime, ampicillin/sulbactam, imipenem,
and amikacin. BPI protein product also increased susceptibility of *P.*
retgeri to ceftazidime and cefuroxime.

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TABLE 25 EFFECTS OF rBPI ₂₁ ± ANTIBIOTICS ON <i>Providencia</i> SPECIES				
Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (µg/mL)		
		With 0 µg/mL rBPI ₂₁	With 8 µg/mL rBPI ₂₁	With 32 µg/mL rBPI ₂₁
19614 (<i>P. reingeri</i>)	Cefoxitin	> 16	8	> 16
	Cefuroxime*	16	8	< 2
	Amikacin	16	16	16
18435 (<i>P. stuartii</i>)	Piperacillin*	> 64	32	16
	Cefazolin**	> 16	> 16	8
	Cefuroxime**	> 16	> 16	8
	Ceftazidime	4	< 2	< 2
	Ampicillin/ Sulbactam	> 16	16	16
	Imipenem*	8	< 4	< 4
	Amikacin	16	16	8

* The Microscan® worksheet did not supply antibiotic susceptibility ranges for this organism, but there was reversal of antibiotic indifference according to NCCLS standards, Publication M7-A3, Table 2 (1993).

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Example 19**EFFECTS OF BPI PROTEIN PRODUCT
AND ANTIBIOTICS *IN VITRO* ON *MORGANELLA MORGANII***

5 The effect of a BPI protein product, rBPI₂₁, on the antibiotic
susceptibility of *Morganella morganii* was evaluated in the Microscan®
antibiotic susceptibility screening assay of Example 11 using the Neg
Combo Type 16 panel plate. The direct growth inhibitory effect of rBPI₂₁
on these strains was also evaluated in the same assay. Assays were
conducted on clinical isolates of *Morganella morganii* (from Baxter
10 Microscan® library, Sacramento, CA).

A summary of the results of the antibiotic screening panels,
reported as MICs (µg/ml) of the antibiotic tested, is shown in Table 26
below. Results are reported for each strain tested, but susceptibility data is
listed for only those antibiotics for which BPI protein product altered
15 susceptibility. The antibiotic susceptibility standards (interpretation of an
MIC as resistant, intermediate or susceptible according to Microscan®'s
NCCLS-derived standards) applicable to the organism tested appear in
Table 13A. These results show that BPI protein product increased the
susceptibility of the tested strains to ampicillin/sulbactam, amikacin and
20 piperacillin.

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TABLE 26 EFFECTS OF rBPI ₂₁ ± ANTIBIOTICS ON <i>Morganella morganii</i>				
Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (µg/mL)		
		With 0 µg/mL rBPI ₂₁	With 8 µg/mL rBPI ₂₁	With 32 µg/mL rBPI ₂₁
F19-004	Ampicillin/ Sulbactam	> 16	16	16
	Amikacin	4	< 2	4
F19-005	Ampicillin/ Sulbactam*	> 16	< 8	< 8
	Amikacin	4	< 2	< 2
F19-006	Piperacillin	> 64	64	64
	Amikacin	< 2	< 2	4

* The Microscan® worksheet did not supply antibiotic susceptibility ranges for this organism, but there was reversal of antibiotic indifference according to NCCLS standards, Publication M7-A3, Table 2 (1993).

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Example 20**EFFECTS OF BPI PROTEIN PRODUCT
AND ANTIBIOTICS *IN VITRO* ON *ACINETOBACTER* SPECIES**

The effect of a BPI protein product, rBPI₂₁, on the antibiotic susceptibility of various *Acinetobacter* species was evaluated in the Microscan® antibiotic susceptibility screening assay of Example 11 using the Neg Combo Type 16 and MIC Plus Type 2 panel plate. The direct growth inhibitory effect of rBPI₂₁ on these strains was also evaluated in the same assay. Assays were conducted on clinical isolates of *Acinetobacter* species (from Baxter Microscan® library, Sacramento, CA). Different production lots of rBPI₂₁ that had been formulated with surfactant or unformulated (without surfactant) were tested, but no difference was seen in results for formulated or unformulated rBPI₂₁.

A summary of the results of the antibiotic screening panels, reported as MICs (µg/ml) of the antibiotic tested, is shown in Tables 27 and 28 below. Results are reported for each strain tested, but susceptibility data is listed for only those antibiotics for which BPI protein product altered susceptibility. The antibiotic susceptibility standards (interpretation of an MIC as resistant, intermediate or susceptible according to Microscan®'s NCCLS-derived standards) applicable to the organism tested appear in Tables 13A and 14A. For interpretation purposes, when antibiotic susceptibility standards were given only for *Enterobacter* or *Pseudomonas*, the standards for *Acinetobacter* were considered to be the same as for *Enterobacter*.

These results show that BPI protein product reversed resistance of *A. anitratus* strains to amoxicillin/K clavulanate, ampicillin/sulbactam, aztreonam, carbenicillin, cefamandole, cefazolin, cefonicid, cefoperazone, cefotaxime, cefotetan, ceftazidime, ceftizoxime, ceftriaxone, chloramphenicol, ciprofloxacin, gentamicin, mezlocillin, netilmicin, ticarcillin, ticarcillin/K clavulanate, and trimethoprim/sulfamethoxazole, and increased susceptibility of some *A. anitratus* strains to amikacin, amoxicillin/K clavulanate, ampicillin,

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ampicillin/sulbactam, azlocillin, aztreonam, carbenicillin, cefamandole,
cefazolin, cefonicid, cefoperazone, cefotaxime, cefotetan, ceftazidime,
ceftazoxime, ceftriaxone, cefuroxime, chloramphenicol,
ciprofloxacin, gentamicin, imipenem, mezlocillin, netilmicin, ofloxacin,
5 piperacillin, ticarcillin, ticarcillin/K clavulanate, tobramycin and
trimethoprim/sulfamethoxazole.

BPI protein product reversed resistance of *A. lwoffii* strains
to aztreonam, cefazolin, cefuroxime, ceftazidime, ceftazoxime,
trimethoprim/sulfamethoxazole and piperacillin, and increased susceptibility
10 of *A. lwoffii* strains to ampicillin, ampicillin/sulbactam, aztreonam,
cefazolin, cefotaxime, ceftazidime, ceftriaxone, cefuroxime,
piperacillin, ticarcillin and trimethoprim/sulfamethoxazole.

These results also show that rBPI₁ at a concentration of 16
μg/ml has direct bactericidal/growth inhibitory effects on some of the tested
15 *Acinetobacter* isolates.

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TABLE 27
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Acinetobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
N011-002 (<i>A. anitratus</i>)	BPI	G	G	G
	Ampicillin	> 16	16	16
	Amikacin	4	<2	<2
N011-003 (<i>A. anitratus</i>)	BPI	G	G	G
	Ampicillin	4	<2	4
	Ticarcillin	16	<8	<8
	Cefazolin**	> 16	> 16	8
	Cefoxitin**	> 16	8	8
	Cefuroxime	8	4	8
N011-070 (<i>A. anitratus</i>)	BPI	G	Reduced	NG
	Ampicillin	8	8	<2
	Trimeth/Sulfa**	>2	2	<0.5
	Cefazolin**	> 16	> 16	<2
	Cefoxitin*	16	16	<2
	Cefuroxime*	16	8	<2
	Ceftazidime	4	<2	<2

TABLE 27
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Acinetobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
N011-071 (<i>A. anitratus</i>)	BPI	G	G	Very reduced
	Ticarcillin*	32	<8	<8
	Aztreonam	>16	16	16
	Piperacillin*	32	16	<8
	Ciprofloxacin**	>2	2	<1
	Ofloxacin*	4	<2	<2
	Ceftazidime	8	8	4
	Ceftriaxone	32	16	16
	Cefotaxime	32	16	16
	Gentamicin**	>6	>6	2
	Tobramycin*	6	4	4
	Amikacin**	>16	16	8
N011-072 (<i>A. anitratus</i>)	BPI	G	Reduced	NG
	Ampicillin*	16	16	<2
	Trimeth/Sulfa**	>2	>2	<0.5
	Cefazolin**	>16	>16	<2
	Cefoxitin*	16	16	<2
	Cefuroxime	8	8	<2
	Gentamicin*	6	4	<1
	Tobramycin	2	<1	<1
	Amikacin	8	4	<2

TABLE 27
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Acinetobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
N012-001 (<i>A. lwoffii</i>)	BPI	G	G	G
	Aztreonam**	> 16	> 16	< 8
	Cefazolin**	> 16	8	< 2
	Cefuroxime**	> 16	4	< 2
	Ceftazidime**	> 16	4	< 2
	Ampicill/Sulbact	> 16	> 16	16
	Cefotaxime	8	< 4	< 4
N012-002 (<i>A. lwoffii</i>)	BPI	G	NG	NG
	Ampicillin	4	< 2	< 2
	Trimeth/Sulfa**	> 2	2	< 0.5
	Aztreonam**	> 16	< 8	< 8
	Cefazolin**	> 16	< 2	< 2
	Cefoxitin**	> 16	< 2	< 2
	Cefuroxime	8	< 2	< 2
	Ceftazidime	4	< 2	< 2

TABLE 27
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Acinetobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
N012-003 (<i>A. lwoffii</i>)	BPI	G	G	NG
	Ampicillin*	> 16	4	< 2
	Trimeth/Sulfa*	2	< 0.5	< 0.5
	Ticarcillin*	64	< 8	< 8
	Aztreonam**	> 16	< 8	< 8
	Piperacillin**	> 64	< 8	< 8
	Cefazolin**	> 16	> 16	< 2
	Cefoxitin**	> 16	8	< 2
	Cefuroxime**	> 16	< 2	< 2
	Ceftazidime*	16	< 2	< 2
	Ceftriaxone*	16	< 4	< 4
	Cefotaxime	8	< 4	< 4
N012-004 (<i>A. lwoffii</i>)	BPI	G	G	G
	Ampicillin	4	< 2	4
	Trimeth/Sulfa*	2	2	< 0.5
	Cefazolin	> 16	16	16
	Cefoxitin*	16	< 2	4
	Cefuroxime	4	< 2	< 2

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TABLE 27 EFFECTS OF rBPI ₂₁ ± ANTIBIOTICS ON <i>Acinetobacter</i> SPECIES				
Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
N012-005 (<i>A. lwoffii</i>)	BPI	G	G	NG
	Ampicillin	4	<2	<2
	Trimeth/Sulfa ^a	2	<0.5	<0.5
	Cefazolin ^{ab}	>16	>16	4
	Cefoxitin ^a	16	8	4
	Cefuroxime	4	4	<2

TABLE 28
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Acinetobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (µg/mL)		
		With 0 µg/mL rBPI ₂₁	With 4 µg/mL rBPI ₂₁	With 16 µg/mL rBPI ₂₁
12292 (<i>A. anitratus</i>)	Ceftizoxime**	> 32	<2	<2
	Ceftazidime**	> 32	32	<1
	Cefotaxime**	> 64	8	<2
	Ceftriaxone**	> 64	<2	<2
	Cefoperazone**	> 32	8	<4
	Cefonicid**	> 16	16	<2
	Cefotetan**	> 32	32	<4
	Netilmicin**	> 16	<2	<2
	Cefamandole**	> 32	<4	<4
	Chloramphenicol**	> 16	<2	<2
	Ticarcillin*	64	32	<16
	Azlocillin	> 64	<64	<64
	Imipenem	4	<0.5	<0.5
	Amp/Sulbactam**	32	4	<1
	Aztreonam**	32	4	<1
	Amox/K Clavulanate*	16	<1	<1
	Ciprofloxacin**	>4	1	<25
	Ticar/K Clavulanate**	64	<16	<16
	Mezlocillin**	> 128	<16	<16
	Carbenicillin**	> 128	<16	<16

TABLE 28
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Acinetobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
12300 (<i>A. anitratus</i>)	Ceftizoxime**	> 32	> 32	< 8
	Ceftazidime	8	8	2
	Cefotaxime**	64	64	< 2
	Ceftriaxone*	32	32	< 2
	Cefoperazone	> 32	> 32	32
	Cefonicid	> 16	> 16	> 16
	Cefotetan	> 32	> 32	> 32
	Netilmicin	> 16	16	16
	Cefamandole**	> 32	> 32	< 4
	Chloramphenicol**	> 16	> 16	8
	Ticarcillin	< 16	< 16	< 16
	Azlocillin	> 64	< 64	< 64
	Imipenem	< 0.5	< 0.5	< 0.5
	Amp/Sulbact	2	2	< 1
	Aztreonam	8	8	4
	Amox/K Clavulanate*	16	16	< 1
	Ciprofloxacin**	> 4	> 4	1
	Ticar/K Clavulanate	< 16	< 16	< 16
	Mezlocillin**	128	128	< 16
	Carbenicillin	< 16	< 16	< 16

TABLE 28
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Acinetobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
12487 <i>(A. anitrans)</i>	Ceftizoxime**	> 32	> 32	< 2
	Ceftazidime**	> 32	2	8
	Cefotaxime**	> 64	64	4
	Ceftriaxone**	> 64	> 64	< 2
	Cefoperazone**	> 32	> 32	< 4
	Cefonicid	> 16	> 16	> 16
	Cefotetan	> 32	> 32	32
	Netilmicin**	> 16	< 2	< 2
	Cefamandole**	> 32	< 4	< 4
	Chloramphenicol**	> 16	8	< 2
	Ticarcillin**	128	32	< 16
	Azlocillin	> 64	< 64	< 64
	Imipenem	2	< 0.5	< 0.5
	Amp/Sulbact*	16	2	< 1
	Aztreonam**	32	16	< 1
	Amox/K Clavulanate	8	< 1	2
	Ciprofloxacin*	2	1	< 0.25
	Ticar/K Clavulanate	32	< 16	< 16
	Mezlocillin**	> 128	32	< 16
	Carbenicillin**	128	< 16	< 16

TABLE 28
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Acinetobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
19687 (<i>A. anitrans</i>)	Ceftizoxime	> 32	> 32	> 32
	Ceftazidime	> 32	> 32	> 32
	Cefotaxime	> 64	> 64	> 64
	Ceftriaxone	> 64	> 64	> 64
	Cefoperazone	> 32	> 32	> 32
	Cefonicid	> 16	> 16	> 16
	Cefotetan	> 32	> 32	> 32
	Netilmicin*	16	8	< 2
	Cefamandole	> 32	> 32	> 32
	Chloramphenicol**	> 16	16	8
	Ticarcillin	> 128	< 16	128
	Azlocillin	> 64	> 64	> 64
	Imipenem	4	2	< 0.5
	Amp/Sulbact**	32	8	2
	Aztreonam	> 32	> 32	> 32
	Amox/K Clavulanate**	32	16	8
	Ciprofloxacin**	> 4	> 4	0.5
	Ticar/K Clavulanate	> 128	128	< 16
	Mezlocillin	> 128	64	32
	Carbenicillin**	> 128	64	< 16

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TABLE 28
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Acinetobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (μg/mL)		
		With 0 μg/mL rBPI ₂₁	With 4 μg/mL rBPI ₂₁	With 16 μg/mL rBPI ₂₁
19693 (<i>A. anitrans</i>)	Ceftizoxime**	> 32	> 32	4
	Ceftazidime	> 32	> 32	> 32
	Cefotaxime	> 64	64	32
	Ceftriaxone	> 64	64	16
	Cefoperazone**	> 32	> 32	8
	Cefonicid**	> 16	> 16	< 2
	Cefotetan**	> 32	> 32	< 4
	Netilmicin	> 16	> 16	> 16
	Cefamandole	> 32	> 32	16
	Chloramphenicol**	> 16	> 16	< 2
	Ticarcillin**	> 128	> 128	< 16
	Azlocillin	> 64	> 64	< 64
	Imipenem	2	2	< 0.5
	Amp/Sulbact**	32	32	2
	Aztreonam**	32	16	< 1
	Amox/K Clavulanate*	16	8	4
	Ciprofloxacin	> 4	> 4	> 4
	Ticar/K Clavulanate**	> 128	32	< 16
	Mezlocillin**	> 128	128	< 16
	Carbenicillin**	> 128	> 128	< 16

TABLE 28
EFFECTS OF rBPI₂₁ ± ANTIBIOTICS ON *Acinetobacter* SPECIES

Microscan Library ID No.	Antibiotic Tested	Minimum Inhibitory Concentration of Antibiotic (µg/mL)		
		With 0 µg/mL rBPI ₂₁	With 4 µg/mL rBPI ₂₁	With 16 µg/mL rBPI ₂₁
19694 (<i>A. anitratus</i>)	Ceftizoxime	> 32	> 32	32
	Ceftazidime*	16	16	8
	Cefotaxime	64	64	64
	Ceftriaxone	64	64	32
	Cefoperazone	> 32	> 32	> 32
	Cefonicid**	> 16	> 16	8
	Cefotetan**	> 32	> 32	8
	Netilmicin**	> 16	> 16	4
	Cefamandole	> 32	> 32	> 32
	Chloramphenicol	> 16	> 16	16
	Ticarcillin	> 128	> 128	32
	Azlocillin	> 64	> 64	> 64
	Imipenem	1	1	< 0.5
	Amp/Sulbact	32	32	16
	Aztreonam	8	8	8
	Amox/K Clavulanate**	32	8	4
	Ciprofloxacin	> 4	> 4	> 4
	Ticar/K Clavulanate**	> 128	< 16	< 16
	Mezlocillin	> 128	> 128	64
	Carbenicillin**	> 128	> 128	< 16

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Example 21**EFFECTS OF BPI PROTEIN PRODUCT AND
ANTIBIOTICS IN VITRO ON *SALMONELLA* AND *SHIGELLA* SPECIES**

Ten clinical isolates of *Salmonella* (F270-001 through -010) and 10
5 clinical isolates of *Shigella* (F321-010 and F325-002 through -010) (all isolates
from Baxter Microscan® library, Sacramento, CA), were evaluated in the
Microscan® antibiotic susceptibility screening assay of Example 11 using the Neg
Breakpoint Combo Type 9 panel plate. Essentially no effect was seen at rBPI₂₁
concentrations of 0, 4 and 16 µg/mL.

Example 22**EFFECTS OF BPI PROTEIN PRODUCT ON
ANTIBIOTIC KILLING CURVES FOR
5 *E. COLI* J5, *E. COLI* O7:K1, *ENTEROBACTER CLOACAE*,
AND *KLEBSIELLA PNEUMONIAE***

The effect of a BPI protein product, rBPI₂₁, on the killing curves of
selected antibiotics was determined for selected organisms. Microscan® panel
plates were prepared for *E. coli* J5, *E. coli* O7:K1, *Enterobacter cloacae*
(Microscan library ID no. 19680) and *Klebsiella pneumoniae* (Microscan library
10 ID no. 16135), according to Example 11. Cell suspensions were added to 25 ml
Pluronic Inoculum Water containing 0 or 16 µg/ml rBPI₂₁. After inoculation, the
panel plates were incubated at 35°C for 24 hours. At 0, 4, 7 and 24 hours after
inoculation, 5 µl samples were removed from each growth control well (containing
culture media without antibiotic) and from each well containing:
15 trimethoprim/sulfamethoxazole, 2 µg/ml ciprofloxacin, 64 µg/ml piperacillin, 32
µg/ml cefotaxime, 6 µg/ml cefuroxime, and 16 µg/ml amikacin. These 5 µl
samples were diluted in sterile water and inoculated onto Trypticase Soy agar
plates (Remel, Lenexa, Kansas). After 48 hours of incubation at 35°C, the plates
were counted and the number of colony forming units of bacteria in the well was
20 calculated.

The results are shown below in Figures 19-25. In all of the figures
the growth, in the presence of antibiotic but without rBPI₂₁, is indicated for: *E.*
coli J5 (a filled square); *E. coli* O7:K1 (a filled diamond); *E. cloacae* (a filled

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triangle); and *K. pneumoniae* (an "X"). Also, in all figures, the growth in the presence of antibiotic with rBPI₂₁ is indicated for *E. coli* J5 (an open square); *E. coli* O7:K1 (an open diamond); *E. cloacae* (an open triangle); and *K. pneumoniae* (a star).

5 Figure 19 shows the kinetic growth curve of organisms with rBPI₂₁ (and without antibiotic) and without rBPI₂₁ (and without antibiotic). In Figure 19, the growth curves for *E. coli* J5 without rBPI₂₁ (filled squares), *E. coli* O7:K1 with rBPI₂₁ (open diamonds), *E. coli* O7:K1 without rBPI₂₁ (filled diamonds), and *K. pneumoniae* without rBPI₂₁ ("X"s) overlap substantially, while the growth curves for *E. coli* J5 with rBPI₂₁ (open squares) and *E. cloacae* with rBPI₂₁ (open triangles) overlap at 0-7 hours but diverge by 24 hours. Figure 19 demonstrates that BPI protein product alone has a bactericidal effect at 0-7 hours on *E. coli* J5 and *K. pneumoniae*, and a bactericidal effect on *E. cloacae* throughout the 24 hour period studied.

15 In Figure 20, the growth curves for *E. coli* J5 with rBPI₂₁ (open squares), *E. cloacae* with rBPI₂₁ (open triangles) and *K. pneumoniae* with rBPI₂₁ (stars) overlap. Figure 20 shows that rBPI₂₁ enhanced the bactericidal effect of trimethoprim/sulfamethoxazole on *E. cloacae* and *K. pneumoniae* at 0-24 hours, and slightly enhanced the antibiotics' effect on *E. coli* J5 and *E. coli* O7:K1.

20 In Figure 21, the growth curves for *E. coli* J5 with rBPI₂₁ (open squares), *E. coli* J5 without rBPI₂₁ (filled squares), *E. coli* O7:K1 with rBPI₂₁ (open diamonds), *E. coli* O7:K1 without rBPI₂₁ (filled diamonds), *E. cloacae* with rBPI₂₁ (open triangles), *E. cloacae* without rBPI₂₁ (filled triangles), and *K. pneumoniae* with rBPI₂₁ (stars) overlap. Figure 21 shows that rBPI₂₁ reversed resistance of *K. pneumoniae* to ciprofloxacin at 0-24 hours; the other organisms were already very susceptible to ciprofloxacin.

25 In Figure 22, the growth curves for *E. coli* O7:K1 without rBPI₂₁ (filled diamonds) and *E. cloacae* without rBPI₂₁ (filled triangles) overlap, while the growth curves for *E. coli* J5 with rBPI₂₁ (open squares), *E. coli* J5 without rBPI₂₁ (filled squares), *E. coli* O7:K1 with rBPI₂₁ (open diamonds), *E. cloacae* with rBPI₂₁ (open triangles), *K. pneumoniae* without rBPI₂₁ ("X"s) and *K. pneumoniae* with rBPI₂₁ (stars) overlap. Figure 22 shows that rBPI₂₁ enhanced the bactericidal

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effect of piperacillin on *E. coli* O7:K1 and *E. cloacae*; the other organisms were already susceptible to the drug.

5 In Figure 23, the growth curves for *E. coli* J5 with rBPI₂₁ (open squares), *E. coli* O7:K1 with rBPI₂₁ (open diamonds), *E. coli* O7:K1 without rBPI₂₁ (filled diamonds), *K. pneumoniae* without rBPI₂₁ ("X"s) and *K. pneumoniae* with rBPI₂₁ (stars) overlap. Figure 23 shows that rBPI₂₁ enhanced the bactericidal effect of cefotaxime on J5 and *E. cloacae*; the other organisms were susceptible to the drug.

10 In Figure 24, all of the curves overlap. Figure 24 shows the effect of rBPI₂₁ and cefuroxime; addition of rBPI₂₁ had no effect on the killing curves because all of the tested species were very susceptible to the antibiotic.

In Figure 25, all of the curves overlap. Figure 25 shows the effect of rBPI₂₁ and amikacin; again, all tested species were very susceptible to the antibiotic.

15

Example 23

EFFECTS OF A VARIETY OF BPI PROTEIN PRODUCTS AND ANTIBIOTICS *IN VITRO* ON REPRESENTATIVE GRAM-NEGATIVE ORGANISMS

20 The effects of a variety of BPI protein products, rBPI₁₁, rBPI₁₃, rBPI₃₀ and rBPI₄₂ dimer on the antibiotic susceptibility of various representative gram-negative organisms was evaluated in the Microscan® antibiotic susceptibility screening assay of Example 11 using the MIC Plus Type 2 panel plate. Assays were conducted on clinical isolates of *Acinetobacter anitratus* and *Enterobacter cloacae* (from Baxter Microscan® library, Sacramento, CA), and on *E. coli* J5-L and O7:K1.

25

The results, reported as MICs (µg/ml) of the antibiotic tested at the various concentrations of BPI protein product indicated, are shown in Tables 29, 30, 31 and 32 below.

TABLE 29
EFFECTS OF BPI PROTEIN PRODUCTS \pm ANTIBIOTICS ON *A. anirralis* (Microscan ID No. 12292)

ANTIBIOTIC TESTED	Minimum Inhibitory Concentration of Antibiotic ($\mu\text{g/ml}$) With:									
	0	4	4	4	4	4	4	4	16	16
	$\mu\text{g/mL}$ BPI	$\mu\text{g/mL}$ BPI ₁₁	$\mu\text{g/mL}$ BPI ₁₁	$\mu\text{g/mL}$ BPI ₁₁	$\mu\text{g/mL}$ BPI ₁₁	$\mu\text{g/mL}$ BPI ₁₁	$\mu\text{g/mL}$ BPI ₁₁	$\mu\text{g/mL}$ BPI ₁₁	$\mu\text{g/mL}$ BPI ₁₁	$\mu\text{g/mL}$ BPI ₁₁
Ceftizoxime	> 32	> 32	> 32	> 32	> 32	> 32	> 32	> 32	> 32	> 32
Ceftazidime	> 32	16	> 32	> 32	> 32	> 32	> 32	> 32	> 32	> 32
Ceftriaxime	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
Ceftriaxone	> 64	8	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
Cefoperazone	> 32	> 32	> 32	> 32	> 32	> 32	> 32	> 32	> 32	> 32
Cefonicid	> 16	> 16	> 16	> 16	> 16	> 16	> 16	> 16	> 16	> 16
Cefotetan	> 32	32	> 32	> 32	> 32	> 32	> 32	> 32	> 32	> 32
Netilmicin	> 16	< 2	> 16	> 16	> 16	> 16	> 16	> 16	> 16	> 16
Cefamandole	> 32	8	> 32	> 32	> 32	> 32	> 32	> 32	> 32	> 32
Chloramphenicol	> 16	< 2	> 16	> 16	> 16	> 16	> 16	> 16	> 16	> 16
Ticarcillin	128	< 16	128	64	64	64	64	64	64	64
Azlocillin	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64	> 64
Imipenem	4	< 0.5	2	2	2	2	2	2	2	2

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TABLE 29 EFFECTS OF BPI PROTEIN PRODUCTS ± ANTIBIOTICS ON <i>A. aviratus</i> (Microscan ID No. 12292)											
ANTIBIOTIC TESTED	Minimum Inhibitory Concentration of Antibiotic (µg/ml) With:										
	0 µg/mL BPI	4 µg/mL BPI ₁₁	4 µg/mL BPI ₁₁	4 µg/mL BPI ₁₁	4 µg/mL BPI ₁₀	4 µg/mL BPI ₄ dimer	16 µg/mL BPI ₁₁	16 µg/mL BPI ₁₁	16 µg/mL BPI ₁₁	16 µg/mL BPI ₁₀	16 µg/mL BPI ₄ dimer
Ampicillin/ Sulhactam	32	4	16	32	32	16	<1	8	16	16	8
Aztreonam	32	16	32	32	32	32	4	32	32	32	32
Amoxicillin/K Clavulanate	16	2	16	16	16	8	4	8	8	8	4
Ciprofloxacin	4	2	4	4	4	4	2	4	4	4	2
Ticarcillin/K Clavulanate	64	<16	64	64	64	32	32	32	32	32	32
Mezlocillin	>128	128	>128	>128	>128	>128	<16	>128	>128	>128	>128
Carbenicillin	128	<16	128	128	128	64	<16	64	64	64	64

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EFFECTS OF BPI PROTEIN PRODUCTS ± ANTIBIOTICS ON <i>E. cloacae</i> (Microscan ID No. 19680)													
ANTIBIOTIC TESTED	Minimum Inhibitory Concentration of Antibiotic (μg/ml) With:												
	0 μg/mL BPI	4 μg/mL BPI ₁₁	4 μg/mL BPI ₁₂	4 μg/mL BPI ₁₃	4 μg/mL BPI ₁₄	4 μg/mL BPI ₁₅	4 μg/mL BPI ₁₆	4 μg/mL BPI ₁₇	4 μg/mL BPI ₁₈	4 μg/mL BPI ₁₉	4 μg/mL BPI ₂₀	4 μg/mL BPI ₂₁	4 μg/mL BPI ₂₂
Cefixoxime	32	32	32	16	32	32	32	32	32	32	32	32	32
Ceftazidime	32	16	16	32	32	32	32	32	32	32	32	32	32
Cefotaxime	64	32	32	32	32	32	32	32	32	32	32	32	32
Ceftriaxone	32	4	32	32	32	32	32	32	32	32	32	32	32
Cefoperazone	32	16	16	16	16	16	16	16	16	16	16	16	16
Cefonicid	>16	>16	>16	>16	>16	>16	>16	>16	>16	>16	>16	>16	>16
Cefotetan	32	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32
Netilmicin	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Cefamandole	>32	>32	32	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32
Chloramphenicol	8	4	4	4	4	4	4	4	4	4	4	4	4
Ticarcillin	>128	64	64	128	128	128	128	128	128	128	128	128	128
Azlocillin	>64	>64	<64	>64	>64	>64	>64	>64	>64	>64	>64	>64	>64
Imipenem	1	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5

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TABLE 30 EFFECTS OF BPI PROTEIN PRODUCTS ± ANTIBIOTICS ON <i>E. cloacae</i> (Microscan ID No. 19680)												
ANTIBIOTIC TESTED	Minimum Inhibitory Concentration of Antibiotic (µg/ml) With:											
	0 µg/mL BPI	4 µg/mL BPI ₁₁	4 µg/mL BPI ₁₃	4 µg/mL BPI ₅₀	4 µg/mL BPI ₄₁ dimer	16 µg/mL BPI ₁₁	16 µg/mL BPI ₁₃	16 µg/mL BPI ₅₀	16 µg/mL BPI ₄₁ dimer	16 µg/mL BPI ₁₁	16 µg/mL BPI ₁₃	16 µg/mL BPI ₅₀
Ampicillin/ Sulbactam	> 32	> 32	> 32	32	> 32	2	2	4	32			
Aztreonam	32	8	4	32	16	< 1	< 1	< 1	16			
Amoxicillin/K Clavulanate	32	32	32	32	32	8	8	16	> 32			
Ciprofloxacin	< .25	< .25	< .25	< .25	< .25	< .25	< .25	< .25	< .25			
Ticarcillin/K Clavulanate	> 128	64	128	128	128	< 16	< 16	< 16	32			
Mezlocillin	< 16	< 16	< 16	< 16	< 16	< 16	< 16	< 16	< 16			
Carbenicillin	> 128	128	64	128	128	< 16	< 16	< 16	128			

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TABLE 31 EFFECTS OF BPI PROTEIN PRODUCTS ± ANTIBIOTICS ON <i>E. coli</i> JS-L												
ANTIBIOTIC TESTED	Minimum Inhibitory Concentration of Antibiotic (µg/ml) With:											
	0 µg/mL BPI	4 µg/mL BPI ₁₁	4 µg/mL BPI ₁₁	4 µg/mL BPI ₁₁	4 µg/mL BPI ₁₀	4 µg/mL BPI ₁₀	4 µg/mL BPI ₁₀	16 µg/mL BPI ₁₁	16 µg/mL BPI ₁₁	16 µg/mL BPI ₁₀	16 µg/mL BPI ₁₀	16 µg/mL BPI ₁₁ dimer
Ceftizoxime	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Ceftazidime	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1
Cefotaxime	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Ceftriaxone	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Cefoperazone	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4
Cefonicid	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Cefotetan	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4
Netilmicin	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Cefamandole	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4
Chloramphenicol	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Ticarcillin	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16
Azlocillin	<64	<64	<64	<64	<64	<64	<64	<64	<64	<64	<64	<64
Imipenem	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5

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TABLE 31 EFFECTS OF BPI PROTEIN PRODUCTS ± ANTIBIOTICS ON <i>E. coli</i> JS-L												
ANTIBIOTIC TESTED	Minimum Inhibitory Concentration of Antibiotic (μg/ml) With:											
	0 μg/mL BPI	4 μg/mL BPI ₁₁	4 μg/mL BPI ₁₃	4 μg/mL BPI ₁₆	4 μg/mL BPI ₁₃ dimer	16 μg/mL BPI ₁₁	16 μg/mL BPI ₁₃	16 μg/mL BPI ₁₆	16 μg/mL BPI ₁₆	16 μg/mL BPI ₁₃ dimer	16 μg/mL BPI ₁₃ dimer	16 μg/mL BPI ₁₃ dimer
Ampicillin/ Sulbactam	2	2	2	2	2	<1	<1	<1	<1	<1	2	2
Aztreonam	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1
Amoxicillin/K Clavulanate	4	<1	2	4	2	<1	<1	<1	2	2	2	2
Ciprofloxacin	<.25	<.25	<.25	<.25	<.25	<.25	<.25	<.25	<.25	<.25	<.25	<.25
Ticarcillin/K Clavulanate	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16
Mezlocillin	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16
Carbenicillin	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16

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TABLE 32
EFFECTS OF BPI PROTEIN PRODUCTS ± ANTIBIOTICS ON *E. coli* O7:K1

ANTIBIOTIC TESTED	Minimum Inhibitory Concentration of Antibiotic (µg/ml) With:									
	0 µg/mL BPI	4 µg/mL BPI ₁₁	4 µg/mL BPI ₁₃	4 µg/mL BPI ₃₀	4 µg/mL BPI ₄₂ dimer	16 µg/mL BPI ₁₁	16 µg/mL BPI ₁₃	16 µg/mL BPI ₃₀	16 µg/mL BPI ₄₂ dimer	16 µg/mL BPI ₄₂ dimer
Ceftazoxime	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Cefazidime	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1
Cefotaxime	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Ceftriaxone	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Cefoperazone	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4
Cefonicid	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Cefotetan	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4
Netilmicin	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Cefamandole	<4	<4	<4	<4	<4	<4	<4	<4	<4	<4
Chloramphenicol	<2	<2	<2	<2	<2	<2	<2	<2	<2	<2
Ticarcillin	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16
Azlocillin	<64	<64	<64	<64	<64	<64	<64	<64	<64	<64
Imipenem	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5	<0.5

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TABLE 32 EFFECTS OF BPI PROTEIN PRODUCTS ± ANTIBIOTICS ON <i>E. coli</i> O7:K1												
ANTIBIOTIC TESTED	Minimum Inhibitory Concentration of Antibiotic (µg/ml) With:											
	0 µg/mL BPI	4 µg/mL BPI ₁₁	4 µg/mL BPI ₁₂	4 µg/mL BPI ₁₃	4 µg/mL BPI ₁₀	4 µg/mL BPI ₁₂ dimer	16 µg/mL BPI ₁₁	16 µg/mL BPI ₁₂	16 µg/mL BPI ₁₀	16 µg/mL BPI ₁₂ dimer	16 µg/mL BPI ₁₁	16 µg/mL BPI ₁₀ dimer
Ampicillin/ Sulbactam	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1
Aztreonam	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1
Amoxicillin/K Clavulanate	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1	<1
Ciprofloxacin	<.25	<.25	<.25	<.25	<.25	<.25	<.25	<.25	<.25	<.25	<.25	<.25
Ticarcillin/K Clavulanate	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16
Mezlocillin	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16
Carbenicillin	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16	<16

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EXAMPLE 24
GRAM-NEGATIVE BACTERICIDAL ACTIVITY OF BPI PEPTIDES

BPI peptides were produced according to co-owned and copending PCT Application No. US94/_____ filed September 15, 1994, which corresponds to U.S. Patent Application Serial No. 08/_____ [Attorney Docket No. 93,1133-H] filed September 15, 1994, and PCT Application No. US94/02465 filed March 11, 1994, which corresponds to U.S. Patent Application Serial No. 08/209,762, filed March 11, 1994, which is a continuation-in-part of U.S. Patent Application Serial No. 08/183,222, filed January 14, 1994, which is a continuation-in-part of U.S. Patent Application Ser. No. 08/093,202 filed July 15, 1993 (for which the corresponding international application is PCT Application No. US94/02401 filed March 11, 1994), which is a continuation-in-part of U.S. Patent Application Ser. No. 08/030,644 filed March 12, 1993, the disclosures of all of which are incorporated herein by reference.

The BPI peptides were screened for bactericidal effects on *E. coli* J5 and *E. Coli* 0111:B4 bacteria in a radial diffusion assay. Specifically, an overnight culture of the bacteria was diluted 1:50 into fresh tryptic soy broth and incubated for 3 hours at 37°C to attain log phase growth of the culture. Bacteria were then pelleted at 3,000 rpm for 5 minutes in a Sorvall RT6000B centrifuge (Sorvall Instruments, Newton, TC). 5 mL of 10 mM sodium phosphate buffer (pH 7.4) was added and the preparation was re-pelleted. The supernatant was decanted and 5 mL of fresh buffer was added, the bacteria were resuspended and their concentration was determined by measurement of absorbance at 590 nm (an Absorbance value of 1.00 at this wavelength equals a concentration of about 1.25×10^9 CFU/mL in suspension). The bacteria were diluted to 4×10^6 CFU/mL in 10 mL of molten underlayer agarose (at approximately 45°C) and inverted repeatedly to mix in 15 mL polypropylene tubes conventionally used for this purpose.

The entire contents of such tubes were then poured into a level square petri dish and distributed evenly by rocking the dish side-to-side. The agarose hardened in less than 30 seconds and had a uniform thickness of about 1 mm. A series of wells were then punched into the hardened agarose using a sterile 3 mm

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punch attached to a vacuum apparatus. The punch was sterilized with 100% alcohol and allowed to air dry prior to use to avoid contaminating the bacterial culture.

5 or 10 μL of each of the BPI peptides were carefully pipetted into each well. As a negative control, dilution buffer (pH 8.3) was added to a separate well, and rBPI₂₂ at concentrations of 5 $\mu\text{g/mL}$ and 1 $\mu\text{g/mL}$ were also added as positive controls. Each plate was incubated at 37°C for 3 hours, and then 10 mL of molten overlayer agarose (at approximately 45°C) was added into the level petri dish, allowed to harden and incubated overnight at 37°C. The next day, a clear zone was seen against the lawn of bacteria in those wells having bactericidal activity. In order to visually enhance this zone, a dilute Coomassie solution (consisting of 0.002% Coomassie Brilliant Blue, 27% methanol, 15% formaldehyde (37% stock solution) and water) was poured over the agar and allowed to stain for 24 hours. The bacterial zones were measured with a micrometer. The assay results for exemplary peptides (BPI.1 through BPI.169) are summarized in Table 33 for the Gram-negative bacteria *E. coli* J5 (rough) and *E. coli* O113 (smooth). The bactericidal activities are expressed as the amount of peptide (pmol/well and $\mu\text{g/well}$) required to generate a 30 mm² bactericidal zone. Additional exemplary BPI peptides include BPI.221 through BPI.281. BPI peptides which retain antibacterial activity are expected to improve the therapeutic effectiveness of antibiotics when concurrently administered therewith. The peptides are screened for such activity in an *in vivo* model or according to *in vitro* tests, including models and tests described herein.

Table 33					
Bactericidal Activity*					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.1	4	-	-	-	-
BPI.2	7	> 2733.5	> 5	-	-
BPI.3	11	696	2.14	-	-
BPI.4	3	-	-	-	-
BPI.5	67	398	1.05	> 1904	> 5
BPI.7	54	175	0.46	> 1890.6	> 5
BPI.8	8	> 3797.1	> 5	-	-
BPI.9	51	479	1.02	> 2345.9	> 5
BPI.10		102	0.41	697	2.76
BPI.11	13	638	1.06	-	-
BPI.12	14	525	1.78	-	-
BPI.13	15	441	0.75	> 2923.9	> 5
BPI.14	2	-	-	-	-
BPI.15	16	> 2733.5	> 5	-	-
BPI.16	17	> 2821.5	> 5	-	-

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Table 33					
Bactericidal Activity ^a					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.17	18	> 2807.2	> 5	-	-
BPI.18	19	> 2757.6	> 5	-	-
BPI.19	20	> 2712.8	> 5	-	-
BPI.20	21	> 2821.5	> 5	-	-
BPI.21	22	> 2917	> 5	-	-
BPI.22	23	> 2821.50	> 5	-	-
BPI.23	24	1330	2.36	> 2821.15	> 5
BPI.24	25	655	1.16	> 2821.50	> 5
BPI.25	26	> 2866.8	> 5	-	-
BPI.26	27	> 2852.1	> 5	-	-
BPI.27	28	> 2797.8	> 5	-	-
BPI.28	29	> 2821.5	> 5	-	-
BPI.29	56	442	1.5	> 1469.2	> 5
BPI.30	52	76	0.23	608	1.84
BPI.31	33	938	1.55	-	-
BPI.32	34	614	1.04	-	-

Table 33					
Bactericidal Activity ^a					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.33	35	575	0.95	-	-
BPI.34	36	916	1.54	-	-
BPI.35	37	263	0.45	-	-
BPI.36	38	1652	2.64	-	-
BPI.37	39	1284	2.14	-	-
BPI.38	40	1698	2.83	-	-
BPI.39	41	316	0.52	-	-
BPI.40	42	1760	2.94	-	-
BPI.41	43	2465	4.03	-	-
BPI.42	44	265	0.44	> 3041.3	> 5
BPI.43	45	729	1.21	> 3024.8	> 5
BPI.44	46	481	0.8	2983	4.93
BPI.45	31	1302	2.23	> 1696.7	> 5
BPI.46	57	186	0.47	> 1811.2	> 5
BPI.47	58	98	0.25	577	1.46
BPI.48	59	42	0.1	254	0.61

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Table 33					
Bactericidal Activity*					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.54	5	-	-	-	-
BPI.55	61	299	0.75	> 1592.2	> 5
BPI.56	47	1387	2.54	-	-
BPI.57	99	514	1.05	-	-
BPI.58	9	1050	2.03	-	-
BPI.59	30	> 2312.3	> 5	-	-
BPI.60	32	> 2136.5	> 5	-	-
BPI.61	48	> 2093.5	> 5	-	-
BPI.63	53	87	0.31	512	1.8
BPI.65 oxidized	10	895	1.82	-	-
BPI.65 reduced	68	1362	2.77	-	-
BPI.66	49	> 3496.7	> 5	-	-
BPI.67	50	> 1901.8	> 5	-	-
BPI.69	60	57	0.21	244	0.88

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Table 33					
Bactericidal Activity ^a					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.70	63	-	-	-	-
BPI.71	64	2297	4.53	-	-
BPI.72	66	> 1911.2	> 5	-	-
BPI.73	62	57	0.11	> 1810.9	> 5
BPI.74	70	732	2.21	> 2148.2	> 5
BPI.75	100	2030.8	4.96	-	-
BPI.76	71	> 3906.5	> 5	-	-
BPI.77	72	455	0.85	-	-
BPI.79	73	> 2282.9	> 5	-	-
BPI.80	74	655	1.24	-	-
BPI.81	75	284	0.52	> 2344.9	> 5
BPI.82	76	171	0.32	> 1197.8	> 5
BPI.83	77	155	0.27	> 2033.5	> 5
BPI.84	78	12	0.02	> 2016.9	> 5
BPI.85	79	227	0.4	> 1881.2	> 5
BPI.86	80	1520	2.58	-	-

Table 33					
Bactericidal Activity ^a					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.87	81	189	0.32	> 1535.8	> 5
BPI.88	82	70.32	0.13	540.15	1
BPI.89	84	229.09	0.43	> 1882.4	> 5
BPI.90	85	83.11	0.16	1763	3.32
BPI.91	86	> 3843.5	> 5	-	-
BPI.92	87	331.8	0.57	-	-
BPI.93	88	212.87	0.76	> 980.3	> 5
BPI.94	89	922.54	1.59	> 922.5	> 5
BPI.95	90	330.88	0.6	> 1397.6	> 5
BPI.96	101	378.33	0.65	> 2048.5	> 5
BPI.97	92	296.58	0.53	-	-
BPI.98	83	> 1626.1	> 5	> 1626.1	> 5
BPI.99	93	722.9	2.99	> 1064.1	> 5
BPI.100	94	407.74	0.73	> 2655	> 5
BPI.101	95	1329.3	4.79	> 1329.3	> 5
BPI.102	96	> 2635.6	> 5	> 2635.6	> 5

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Table 33					
Bactericidal Activity*					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.103	102	165.18	0.31	415.19	0.78
BPI.104	103	385.85	0.64	1376.42	2.30
BPI.105	104	65.35	0.12	206.98	0.39
BPI.106	105	427.12	0.72	> 3413.80	> 5
BPI.107	106	384.67	0.68	> 2795.70	> 5
BPI.108	107	661.05	1.17	> 3219.02	> 5
BPI.109	108	306.80	0.54	> 2822.90	> 5
BPI.110	109	812.33	1.44	> 2950.15	> 5
BPI.111	110	959.00	1.71	> 2808.69	> 5
BPI.112	111	1485.92	2.84	-	-
BPI.113	112	270.66	0.50	> 2950.15	> 5
BPI.114	113	2329.68	3.10	-	-
BPI.116	114	73.82	0.13	> 2788.19	> 5
BPI.119	115	106.70	0.20	536.44	1.02
BPI.120	116	-	-	-	-
BPI.121	117	154.35	0.3	1856.40	3.55

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Table 33					
Bactericidal Activity ^a					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.122	118	179.89	0.36	2123.57	4.2
BPI.123	119	247.20	0.43	> 2865.02	> 5
BPI.124	120	91.23	0.17	> 2580.12	> 5
BPI.125	121	428.85	0.75	> 3149.74	> 5
BPI.126	122	1979.97	3.39	-	-
BPI.127	123	406.01	0.68	-	-
BPI.128	124	2271.14	3.80	-	-
BPI.129	125	1685.10	2.90	-	-
BPI.130	126	325.75	0.68	> 2903.34	> 5
BPI.131	127	1438.21	2.48	-	-
BPI.132	128	> 2988.50	> 5	-	-
BPI.133	129	2316.59	3.91	-	-
BPI.134	130	162.5	0.30	580.11	1.05
BPI.135	131	1052.02	1.74	3321.69	> 5
BPI.136	132	> 3030.74	> 5	-	-
BPI.137	133	N.T.	N.T.	N.T.	N.T.

Table 33					
Bactericidal Activity*					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.138	134	64.57	0.11	995.40	1.74
BPI.139	135	1261.37	2.13	3793.91	> 5
BPI.140	136	84.76	0.26	605.34	1.89
BPI.141	137	> 2809.51	> 5	-	-
BPI.142	138	922.21	1.76	-	-
BPI.143	139	> 2838.99	> 5	-	-
BPI.144	140	510.02	0.86	-	-
BPI.145	141	N.T.	N.T.	N.T.	N.T.
BPI.146	142	-	-	-	-
BPI.147	143	> 2558.17	> 5	-	-
BPI.148	144	> 2805.45	> 5	-	-
BPI.149	147	44.00	0.57	391.00	5.00
BPI.150	148	220.00	0.58	> 2380.67	> 5
BPI.151		N.T.	N.T.	N.T.	N.T.
BPI.152		N.T.	N.T.	N.T.	N.T.
BPI.153	149	N.T.	N.T.	N.T.	N.T.

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Table 33					
Bactericidal Activity*					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.154	150	197.00	0.55	2977.76	> 5
BPI.155	151	> 1795.66	> 5	> 1795.66	> 5
BPI.156	152	N.T.	N.T.	N.T.	N.T.
BPI.157	153	N.T.	N.T.	N.T.	N.T.
BPI.158	154	N.T.	N.T.	N.T.	N.T.
BPI.159	155	765.43	2.41	> 1589.88	> 5
BPI.160	156	288.78	0.81	> 1781.59	> 5
BPI.161	157	1201.79	2.00	-	-
BPI.162	158	N.T.	N.T.	N.T.	N.T.
BPI.163	159	N.T.	N.T.	N.T.	N.T.
BPI.164	160	N.T.	N.T.	N.T.	N.T.
BPI.165	161	N.T.	N.T.	N.T.	N.T.
BPI.166	162	514.00	0.83	> 3078.72	> 5
BPI.167	163	> 4585.73	> 5	-	-
BPI.168	164	1460.98	2.87	> 1948.48	> 5

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Table 33					
Bactericidal Activity ^a					
BPI Peptide	Sequence ID No.	<i>E. coli</i> J5		<i>E. coli</i> O111:B4	
		(pmol/well)	(μ g/well)	(pmol/well)	(μ g/well)
BPI.169	165	4893.83	> 5	> 4974.43	> 5
BPI.170	227	3693.06	> 5	-	-
MAP.1 [*]		106	0.82	552.79	4.27
MAP.2 ^{**}		> 690.9	> 5	> 690.9	> 5

a Amount added to well to achieve a 30 mm² hole as determined by PROBIT analysis as described in Examples 15 and 16.

b No detectable activity up to 5 μ g/well.

c N.T. = not tested

^{*} MAP.1 = β -Ala-N α ,NE-([N α ,NE(BPI.2)Lys]Lys

^{**} MAP.2 = β -Ala-N α ,NE-([N α ,NE(BPI.13)Lys]Lys

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EXAMPLE 25
EFFECTS OF CONCURRENT ADMINISTRATION
OF BPI PROTEIN PRODUCT
AND TETRACYCLINE OR GENTAMICIN ON *E. COLI* 0111:B4

5 Additional MIC assays were performed to determine the sensitivity of *E. coli* 0111:B4 (as described in Example 1) to the effects of BPI protein products concurrently administered with the antibiotic tetracycline or with the antibiotic gentamicin.

10 For these experiments, organisms were grown overnight at 37°C in 5 mL of Mueller-Hinton broth. This overnight culture was diluted 1:50 into 5 mL of fresh broth and incubated for an additional 3 hours at 37°C to attain log-phase growth. Bacteria were pelleted for 5 minutes at 1500 x g and resuspended in fresh broth to give a final concentration of 2×10^6 cells per mL.

15 rBPI₂₂ and antibiotic (either tetracycline or gentamicin) were diluted such that 100 µl bacterial suspension, 50 µl antibiotic and 50 µl diluted rBPI₂₂, gave concentrations in serial dilutions from 10 µg/mL tetracycline or 2.5 µg/mL gentamicin and from 30 µg/mL rBPI₂₂ with a fixed concentration of 10^6 cells/mL. Incubation was carried out in flat bottom 96 well microtiter plates for 18 hours at 37°C, and the plates were read in an automatic plate reader (Titretek Multiscan) at 20 590 nm.

A 4-6 fold decrease in absorbance was observed with tetracycline or gentamicin with certain concentrations of BPI protein product. The MIC of tetracycline without rBPI₂₂, of 10 µg/mL and was reduced by rBPI₂₂ to 5 µg/mL. The MIC of gentamicin without rBPI₂₂, was 0.6 µg/mL and was reduced by rBPI₂₂ 25 to 0.3 µg/mL.

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A partial summary of the data described in the foregoing examples, grouped by general classes of antibiotics, appears below in Table 34 and is displayed as the effects of BPI protein products on the therapeutic effectiveness of antibiotics for various gram-negative organisms.

TABLE 34	
ANTIBIOTIC CLASS	EFFECTS OF BPI PROTEIN PRODUCT WHEN CONCURRENTLY ADMINISTERED WITH ANTIBIOTICS WITHIN CLASS
β -lactams: penicillins and cephalosporins	<p>reversed resistance of <i>Pseudomonas aeruginosa</i>, other <i>Pseudomonas</i>, <i>Xanthomonas</i>, <i>E. coli</i>, <i>Citrobacter</i>, <i>Klebsiella</i>, <i>Enterobacter</i>, <i>Serratia</i>, <i>Providencia</i>, <i>Acinetobacter</i></p> <p>increased susceptibility of <i>Pseudomonas aeruginosa</i>, other <i>Pseudomonas</i>, <i>E. coli</i>, <i>Citrobacter</i>, <i>Klebsiella</i>, <i>Enterobacter</i>, <i>Serratia</i>, <i>Proteus</i>, <i>Providencia</i>, <i>Morganella</i>, <i>Acinetobacter</i></p> <p>enhanced early bactericidal effect for <i>E. coli</i>, <i>Enterobacter</i> in killing curves</p> <p><i>in vivo</i> synergy shown for treatment of <i>E. coli</i> infection in mouse and rabbit models</p>
β -lactams other than penicillins and cephalosporins: aztreonam and imipenem	<p>aztreonam: reversed resistance of <i>Pseudomonas aeruginosa</i>, <i>Acinetobacter</i> increased susceptibility of <i>Acinetobacter</i></p> <p>increased susceptibility of <i>Pseudomonas aeruginosa</i>, <i>Citrobacter</i>, <i>Enterobacter</i></p> <p>imipenem: increased susceptibility of <i>Pseudomonas aeruginosa</i>, <i>Proteus</i>, <i>Providencia</i>, <i>Acinetobacter</i></p>

TABLE 34

ANTIBIOTIC CLASS	EFFECTS OF BPI PROTEIN PRODUCT WHEN CONCURRENTLY ADMINISTERED WITH ANTIBIOTICS WITHIN CLASS
aminoglycosides	<p>reversed resistance of <i>Pseudomonas aeruginosa</i>, <i>Xanthomonas</i>, <i>Acinetobacter</i></p> <p>increased susceptibility of <i>Pseudomonas aeruginosa</i>, other <i>Pseudomonas</i>, <i>E. coli</i>, <i>Citrobacter</i>, <i>Serratia</i>, <i>Proteus</i>, <i>Providencia</i>, <i>Morganella</i>, <i>Acinetobacter</i></p> <p>checkerboard synergy (FIC < 0.5) for <i>E. coli</i>, <i>Salmonella</i>, <i>Klebsiella</i>, <i>Edwardsiella</i>; greater than additive interaction (FIC ≤ 1.0) for <i>Pseudomonas aeruginosa</i>, <i>Enterobacter</i></p> <p><i>in vivo</i> synergy shown for treatment of <i>E. coli</i> infection in mouse model</p>
sulfonamides and trimethoprim	<p>reversed resistance of <i>Pseudomonas aeruginosa</i>, <i>Xanthomonas</i>, <i>Klebsiella</i>, <i>Acinetobacter</i></p> <p>increased susceptibility of <i>Enterobacter</i>, <i>Proteus</i>, <i>Acinetobacter</i></p> <p>enhanced early bactericidal effect for <i>Klebsiella</i>, <i>Enterobacter</i> in killing curves; slight early enhancement for <i>E. coli</i></p>
fluoroquinolones and quinolones	<p>reversed resistance of <i>Pseudomonas aeruginosa</i>, <i>Acinetobacter</i></p> <p>increased susceptibility of <i>Pseudomonas aeruginosa</i>, <i>Xanthomonas</i>, <i>Enterobacter</i>, <i>Acinetobacter</i></p> <p>reversed resistance of <i>Klebsiella</i> in antibiotic killing curves</p>
polymyxins	checkerboard synergy (FIC < 0.5) for <i>Pseudomonas aeruginosa</i> , <i>E. coli</i> , <i>Providencia</i> ; additive interaction (FIC ≤ 1.0) for <i>Enterobacter</i>
chloramphenicol	reversed resistance and increased susceptibility for <i>Acinetobacter</i>

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Numerous modifications and variations in the practice of the invention are expected to occur to those skilled in the art upon consideration of the foregoing description of the presently preferred embodiments thereof. Consequently, the only limitations which should be placed upon the scope of the present invention are those which appear in the appended claims.

5

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANTS: Cohen, Jonathan
Kung, Ada H.C.
Lambert, Jr., Lewis H.
- (ii) TITLE OF INVENTION: Method for Treating Gram-Negative Bacterial Infection by Administration of Bactericidal/Permeability-Increasing (BPI) Protein Product and Antibiotic
- (iii) NUMBER OF SEQUENCES: 227
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 - (B) COMPUTER: IBM PC compatible
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 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
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 - (B) FILING DATE: 22-SEP-1993
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(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 29 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "Domain I"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

142

Ala Ser Gln Gln Gly Thr Ala Ala Leu Gln Lys Glu Leu Lys Arg
 1 5 10 15

Ile Lys Ile Pro Asp Tyr Ser Asp Ser Phe Lys Ile Lys His
 20 25

(2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 30 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.14"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Gly Thr Ala Ala Leu Gln Lys Glu Leu Lys Arg Ile Lys Ile Pro
 1 5 10 15

Asp Tyr Ser Asp Ser Phe Lys Ile Lys His Leu Gly Lys Gly His
 20 25 30

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 22 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.4"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Leu Gln Lys Glu Leu Lys Arg Ile Lys Ile Pro Asp Tyr Ser Asp
 1 5 10 15

Ser Phe Lys Ile Lys His Leu
 20

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 15 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.1"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Gln Gln Gly Thr Ala Ala Leu Gln Lys Glu Leu Lys Arg Ile Lys
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:5:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 15 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.54"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Gly Thr Ala Ala Leu Gln Lys Glu Leu Lys Arg Ile Lys Ile Pro
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:6:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 35 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "Domain II"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

Ser Ser Gln Ile Ser Met Val Pro Asn Val Gly Leu Lys Phe Ser
 1 5 10 15

Ile Ser Asn Ala Asn Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys
 20 25 30

Arg Phe Leu Lys
 35

(2) INFORMATION FOR SEQ ID NO:7:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 15 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.2"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:8:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 10 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

144

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.8"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:9:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 16 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.58"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

Cys Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu
1 5 10 15
Lys

(2) INFORMATION FOR SEQ ID NO:10:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 17 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.65 oxidized"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

Cys Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu
1 5 10 15
Lys Cys

(2) INFORMATION FOR SEQ ID NO:11:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 27 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.3"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

145

Asn Val Gly Leu Lys Phe Ser Ile Ser Asn Ala Asn Ile Lys Ile
 1 5 10 15
 Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
 20 25

(2) INFORMATION FOR SEQ ID NO:12:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 28 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "Domain III"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

Val His Val His Ile Ser Lys Ser Lys Val Gly Trp Leu Ile Gln
 1 5 10 15
 Leu Phe His Lys Lys Ile Glu Ser Ala Leu Arg Asn Lys
 20 25

(2) INFORMATION FOR SEQ ID NO:13:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 13 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.11"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

Lys Ser Lys Val Trp Leu Ile Gln Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:14:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 29 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.12"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Ser Val His Val His Ile Ser Lys Ser Lys Val Gly Trp Leu Ile
 1 5 10 15
 Gln Leu Phe His Lys Lys Ile Glu Ser Ala Leu Arg Asn Lys
 20 25

(2) INFORMATION FOR SEQ ID NO:15:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.13"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:16:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.15"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

Ala Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:17:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.16"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

Ile Ala Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:18:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.17"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

Ile Lys Ala Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:19:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.18"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

Ile Lys Ile Ala Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:20:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.19"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

Ile Lys Ile Ser Ala Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:21:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.20"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

Ile Lys Ile Ser Gly Ala Trp Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:22:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

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- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.21"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:
- Ile Lys Ile Ser Gly Lys Ala Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15
- (2) INFORMATION FOR SEQ ID NO:23:
- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.22"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:
- Ile Lys Ile Ser Gly Lys Trp Ala Ala Gln Lys Arg Phe Leu Lys
1 5 10 15
- (2) INFORMATION FOR SEQ ID NO:24:
- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.23"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:
- Ile Lys Ile Ser Gly Lys Trp Lys Ala Ala Lys Arg Phe Leu Lys
1 5 10 15
- (2) INFORMATION FOR SEQ ID NO:25:
- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.24"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:
- Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Ala Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:26:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.25"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Ala Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:27:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.26"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Ala Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:28:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.27"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Ala Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:29:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc_feature

150

(D) OTHER INFORMATION: "BPI.28"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu Ala
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:30:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.59"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:

Ile Lys Ile Ser Gly Ala Trp Ala Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:31:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.45"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Ala Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:32:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.60"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:

Ile Ala Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu Ala
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:33:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid

151

- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.31"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:
Ala Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10
- (2) INFORMATION FOR SEQ ID NO:34:
- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.32"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:
Lys Ala Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10
- (2) INFORMATION FOR SEQ ID NO:35:
- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.33"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:
Lys Ser Ala Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10
- (2) INFORMATION FOR SEQ ID NO:36:
- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.34"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:
Lys Ser Lys Ala Gly Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:37:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.35"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:

Lys Ser Lys Val Ala Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:38:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.36"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:39:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.37"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:

Lys Ser Lys Val Gly Trp Ala Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:40:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.38"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:

Lys Ser Lys Val Gly Trp Leu Ala Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:41:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.39"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:

Lys Ser Lys Val Gly Trp Leu Ile Ala Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:42:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.40"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:42:

Lys Ser Lys Val Gly Trp Leu Ile Gln Ala Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:43:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.41"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Ala His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:44:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

154

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.42"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe Ala Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:45:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.43"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Ala Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:46:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.44"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Ala
1 5 10

(2) INFORMATION FOR SEQ ID NO:47:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.56"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Lys Gln Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:48:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids

155

(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.61"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:48:

Ile Lys Ile Ser Gly Lys Phe Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:49:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.66"

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 7
(D) OTHER INFORMATION: /label= D-Trp
/note= "The amino acid at position 7 is
D-tryptophan"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:49:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:50:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.67"

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6..8
(D) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 7 is
beta-1-naphthyl-substituted"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:50:

Ile Lys Ile Ser Gly Lys Ala Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:51:

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.9"

Lys Arg Phe Leu Lys Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 24 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.30"

Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys Lys Ser Lys Val Gly
1 5 10 15

Trp Leu Ile Gln Leu Phe His Lys Lys
20

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 29 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.63"

116 Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
1 5 10 15

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
20 25

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

157

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.7"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:54:

Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys Lys Trp Lys Ala Gln
 1 5 10 15
 Lys Arg Phe Leu Lys
 20

(2) INFORMATION FOR SEQ ID NO:55:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 25 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.10.1"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:55:

Lys Arg Phe Leu Lys Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
 1 5 10 15
 Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
 20 25

(2) INFORMATION FOR SEQ ID NO:56:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.29"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:56:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys Lys
 1 5 10 15
 Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
 20 25

(2) INFORMATION FOR SEQ ID NO:57:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.46"

158

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:57:

Lys Trp Lys Ala Ala Ala Arg Phe Leu Lys Lys Trp Lys Ala Gln
1 5 10 15
Lys Arg Phe Leu Lys
20

(2) INFORMATION FOR SEQ ID NO:58:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.47"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:58:

Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys Lys Trp Lys Ala Ala
1 5 10 15
Ala Arg Phe Leu Lys
20

(2) INFORMATION FOR SEQ ID NO:59:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.48"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:59:

Lys Trp Lys Ala Ala Ala Arg Phe Leu Lys Lys Trp Lys Ala Ala
1 5 10 15
Ala Arg Phe Leu Lys
20

(2) INFORMATION FOR SEQ ID NO:60:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 30 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.69"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:60:

Lys Trp Lys Ala Ala Ala Arg Phe Leu Lys Lys Trp Lys Ala Ala
 1 5 10 15
 Ala Arg Phe Leu Lys Lys Trp Lys Ala Ala Ala Arg Phe Leu Lys
 20 25 30

(2) INFORMATION FOR SEQ ID NO:61:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 21 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.55"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:61:

Gly Trp Leu Ile Gln Leu Phe His Lys Lys Ile Glu Ser Ala Leu
 1 5 10 15
 Arg Asn Lys Met Asn Ser
 20

(2) INFORMATION FOR SEQ ID NO:62:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 15 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.73"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:62:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Phe Arg Phe Leu Lys
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:63:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 15 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.70"

- (ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 8..10
 (D) OTHER INFORMATION: /label= Substituted-Ala
 /note= "The alanine at position 7 is
 beta-3-pyridyl-substituted"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:63:

160

Ile Lys Ile Ser Gly Lys Lys Ala Gln Lys Arg Phe Leu Lys
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:64:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.71"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 13..15
- (D) OTHER INFORMATION: /label= Substituted-Ala
 /note= "The alanine at position 13 is
 beta-3-pyridyl-substituted"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:64:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Ala Leu Lys
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:65:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 26 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.10.2"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:65:

Gln Lys Arg Phe Leu Lys Lys Trp Lys Ala Gln Lys Arg Phe Leu
 1 5 10 15
 Lys Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
 20 25

(2) INFORMATION FOR SEQ ID NO:66:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 17 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.72"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 1..3
- (D) OTHER INFORMATION: /label= D-alanine

161

/note= "The position 1 and position 2 alanine residues are both D-alanine"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:66:

Ala Ala Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe
1 5 10 15
Leu Lys

(2) INFORMATION FOR SEQ ID NO:67:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 22 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.5"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:67:

Val His Val His Ile Ser Lys Ser Lys Val Gly Trp Leu Ile Gln
1 5 10 15
Leu Phe His Lys Lys Ile Glu
20

(2) INFORMATION FOR SEQ ID NO:68:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 17 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.65 reduced"

- (ix) FEATURE:
(A) NAME/KEY: Disulfide-bond
(B) LOCATION: 1..17

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:68:

Cys Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe Leu
1 5 10 15
Lys Cys

(2) INFORMATION FOR SEQ ID NO:69:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 487 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

- (ix) FEATURE:
(A) NAME/KEY: misc_feature

162

(D) OTHER INFORMATION: "rSPI"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:69:

Met Arg Glu Asn Met Ala Arg Gly Pro Cys Asn Ala Pro Arg Trp Val
 -31 -30 -25 -20
 Ser Leu Met Val Leu Val Ala Ile Gly Thr Ala Val Thr Ala Ala Val
 -15 -10 -5 1
 Asn Pro Gly Val Val Val Arg Ile Ser Gln Lys Gly Leu Asp Tyr Ala
 5 10 15
 Ser Gln Gln Gly Thr Ala Ala Leu Gln Lys Glu Leu Lys Arg Ile Lys
 20 25 30
 Ile Pro Asp Tyr Ser Asp Ser Phe Lys Ile Lys His Leu Gly Lys Gly
 35 40 45
 His Tyr Ser Phe Tyr Ser Met Asp Ile Arg Glu Phe Gln Leu Pro Ser
 50 55 60 65
 Ser Gln Ile Ser Met Val Pro Asn Val Gly Leu Lys Phe Ser Ile Ser
 70 75 80
 Asn Ala Asn Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe
 85 90 95
 Leu Lys Met Ser Gly Asn Phe Asp Leu Ser Ile Glu Gly Met Ser Ile
 100 105 110
 Ser Ala Asp Leu Lys Leu Gly Ser Asn Pro Thr Ser Gly Lys Pro Thr
 115 120 125
 Ile Thr Cys Ser Ser Cys Ser Ser His Ile Asn Ser Val His Val His
 130 135 140 145
 Ile Ser Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
 150 155 160
 Ile Glu Ser Ala Leu Arg Asn Lys Met Asn Ser Gln Val Cys Glu Lys
 165 170 175
 Val Thr Asn Ser Val Ser Ser Lys Leu Gln Pro Tyr Phe Gln Thr Leu
 180 185 190
 Pro Val Met Thr Lys Ile Asp Ser Val Ala Gly Ile Asn Tyr Gly Leu
 195 200 205
 Val Ala Pro Pro Ala Thr Thr Ala Glu Thr Leu Asp Val Gln Met Lys
 210 215 220 225
 Gly Glu Phe Tyr Ser Glu Asn His His Asn Pro Pro Phe Ala Pro
 230 235 240
 Pro Val Met Glu Phe Pro Ala Ala His Asp Arg Met Val Tyr Leu Gly
 245 250 255
 Leu Ser Asp Tyr Phe Phe Asn Thr Ala Gly Leu Val Tyr Gln Glu Ala
 260 265 270
 Gly Val Leu Lys Met Thr Leu Arg Asp Asp Met Ile Pro Lys Glu Ser
 275 280 285

Lys Phe Arg Leu Thr Thr Lys Phe Phe Gly Thr Phe Leu Pro Glu Val
 290 295 300 305
 Ala Lys Lys Phe Pro Asn Met Lys Ile Gln Ile His Val Ser Ala Ser
 310 315 320
 Thr Pro Pro His Leu Ser Val Gln Pro Thr Gly Leu Thr Phe Tyr Pro
 325 330 335
 Ala Val Asp Val Gln Ala Phe Ala Val Leu Pro Asn Ser Ser Leu Ala
 340 345 350
 Ser Leu Phe Leu Ile Gly Met His Thr Thr Gly Ser Met Glu Val Ser
 355 360 365
 Ala Glu Ser Asn Arg Leu Val Gly Glu Leu Lys Leu Asp Arg Leu Leu
 370 375 380 385
 Leu Glu Leu Lys His Ser Asn Ile Gly Pro Phe Pro Val Glu Leu Leu
 390 395 400
 Gln Asp Ile Met Asn Tyr Ile Val Pro Ile Leu Val Leu Pro Arg Val
 405 410 415
 Asn Glu Lys Leu Gln Lys Gly Phe Pro Leu Pro Thr Pro Ala Arg Val
 420 425 430
 Gln Leu Tyr Asn Val Val Leu Gln Pro His Gln Asn Phe Leu Leu Phe
 435 440 445
 Gly Ala Asp Val Val Tyr Lys
 450 455

(2) INFORMATION FOR SEQ ID NO:70:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 24 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.74"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:70:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys Lys
 1 5 10 15
 Trp Lys Ala Gln Lys Arg Phe Leu Lys
 20

(2) INFORMATION FOR SEQ ID NO:71:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 15 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.76"

164

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 10..12
- (D) OTHER INFORMATION: /label= D-Phe
/note= "The amino acid at position 11 is
D-phenylalanine"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:71:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Phe Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:72:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.77"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:72:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Trp Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:73:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.79"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:73:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Lys Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:74:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.80"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 10..12

165

(D) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 11 is
beta-1-naphthyl-substituted"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:74:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Ala Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:75:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.81"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:75:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Phe Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:76:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.82"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:76:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Trp His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:77:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.83"

(ix) FEATURE:

(A) NAME/KEY: Modified-site
(B) LOCATION: 10..12
(D) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 6 is
beta-1-naphthyl-substituted"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:77:

166

Lys Ser Lys Val Gly Ala Lys Ile Gln Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:78:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.84"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6..8
- (D) OTHER INFORMATION: /label= Substituted-Ala
 /note= "The alanine at position 7 is
 beta-1-naphthyl-substituted"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:78:

Ile Lys Ile Ser Gly Lys Ala Lys Ala Gln Phe Arg Phe Leu Lys
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:79:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.85"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:79:

Lys Ser Lys Val Leu Trp Leu Ile Gln Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:80:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.86"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:80:

Lys Ser Lys Val Gly Trp Leu Ile Leu Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:81:

167

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 14 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.87"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:81:
- Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe Leu Lys Lys
 1 5 10
- (2) INFORMATION FOR SEQ ID NO:82:
- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 15 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.88"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:82:
- Ile Lys Ile Ser Gly Lys Trp Lys Ala Phe Phe Arg Phe Leu Lys
 1 5 10 15
- (2) INFORMATION FOR SEQ ID NO:83:
- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 24 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.98"
- (ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 2
 (D) OTHER INFORMATION: /label= Substituted-Trp
 /note= "The alanine at position 2 is
 beta-1-naphthyl-substituted"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:83:
- Lys Trp Lys Ala Gln Phe Arg Phe Leu Lys Lys Ser Lys Val Gly
 1 5 10 15
- Trp Leu Ile Phe Leu Phe His Lys Lys
 20

(2) INFORMATION FOR SEQ ID NO:84:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.89"

- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6..8
(D) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 7 is
beta-1-naphthyl-substituted"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:84:

Ile Lys Ile Ser Gly Lys Ala Lys Ala Phe Lys Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:85:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.90"

- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6..8
(D) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 7 is
beta-1-naphthyl-substituted"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:85:

Ile Lys Ile Ser Gly Lys Ala Lys Ala Phe Phe Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:86:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.91"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:86:

169

Lys Ser Lys Val Gly Trp Leu Ile Phe Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:87:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 14 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.92"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:87:

Lys Ser Lys Val Gly Trp Leu Ile Lys Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:88:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 29 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.93"

- (ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 6..8
 (D) OTHER INFORMATION: /label= Substituted-Ala
 /note= "The alanine at position 7 is
 beta-1-naphthyl-substituted"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:88:

Ile Lys Ile Ser Gly Lys Ala Lys Ala Gln Phe Arg Phe Leu Lys
 1 5 10 15

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
 20 25

(2) INFORMATION FOR SEQ ID NO:89:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 14 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.94"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:89:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe Phe Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:90:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.95"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:90:

Lys Ser Lys Val Phe Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:91:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.96"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:91:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Phe
1 5 10

(2) INFORMATION FOR SEQ ID NO:92:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.97"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:92:

Lys Ser Lys Val Lys Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:93:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 30 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.99"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:93:

Lys Trp Lys Ala Gln Trp Arg Phe Leu Lys Lys Trp Lys Ala Gln
1 5 10 15
Trp Arg Phe Leu Lys Lys Trp Lys Ala Gln Trp Arg Phe Leu Lys
20 25 30

(2) INFORMATION FOR SEQ ID NO:94:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc_feature
(D) OTHER INFORMATION: "BPI.100"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:94:

Lys Ser Lys Val Lys Trp Leu Ile Lys Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:95:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 28 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc_feature
(D) OTHER INFORMATION: "BPI.101"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:95:

Lys Ser Lys Val Lys Trp Leu Ile Lys Leu Phe Phe Lys Phe Lys
1 5 10 15
Ser Lys Val Lys Trp Leu Ile Lys Leu Phe Phe Lys Phe
20 25

(2) INFORMATION FOR SEQ ID NO:96:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 24 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc_feature
(D) OTHER INFORMATION: "BPI.102"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:96:

Lys Trp Lys Ala Gln Phe Arg Phe Leu Lys Lys Ser Lys Val Gly
1 5 10 15

Trp Leu Ile Leu Leu Phe His Lys Lys
20

(2) INFORMATION FOR SEQ ID NO:97:

(1) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 1443 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..1443

(ix) FEATURE:

- (A) NAME/KEY: mat_peptide
- (B) LOCATION: 76..1443

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "rLBP"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:97:

ATG GGG GCC TTG GCC AGA GCC CTG CCG TCC ATA CTG CTG GCA TTG CTG	48
Met Gly Ala Leu Ala Arg Ala Leu Pro Ser Ile Leu Leu Ala Leu Leu	
-25 -20 -15 -10	
CTT ACG TCC ACC CCA GAG GCT CTG GGT GCC AAC CCC GGC TTG GTC GCC	96
Leu Thr Ser Thr Pro Glu Ala Leu Gly Ala Asn Pro Gly Leu Val Ala	
-5 1 5	
AGG ATC ACC GAC AAG GGA CTG CAG TAT GCG GCC CAG GAG GCG CTA TTG	144
Arg Ile Thr Asp Lys Gly Leu Gln Tyr Ala Ala Gln Glu Gly Leu Leu	
10 15 20	
GCT CTG CAG AGT GAG CTG CTC AGG ATC ACG CTG CCT GAC TTC ACC GGG	192
Ala Leu Gln Ser Glu Leu Leu Arg Ile Thr Leu Pro Asp Phe Thr Gly	
25 30 35	
GAC TTG AGG ATC CCC CAC GTC GCC CGT GGG CCG TAT GAG TTC CAC AGC	240
Asp Leu Arg Ile Pro His Val Gly Arg Gly Arg Tyr Glu Phe His Ser	
40 45 50 55	
CTG AAC ATC CAC AGC TGT GAG CTG CTT CAC TCT GCG CTC AGG CCT GTC	288
Leu Asn Ile His Ser Cys Glu Leu Leu His Ser Ala Leu Arg Pro Val	
60 65 70	
CCT GGC CAG GGC CTG AGT CTC AGC ATC TCC CAC TCC TCC ATC CGG GTC	336
Pro Gly Gln Gly Leu Ser Leu Ser Ile Ser Asp Ser Ser Ile Arg Val	
75 80 85	
CAG GGC AGG TCG AAG GTG CCC AAG TCA TTC TTC AAA CTA CAG GGC TCC	384
Gln Gly Arg Trp Lys Val Arg Lys Ser Phe Phe Lys Leu Gln Gly Ser	
90 95 100	
TIT GAT GTC AGT GTC AAG GGC ATC AGC ATT TCG GTC AAC CTC CTG TTG	432
Phe Asp Val Ser Val Lys Gly Ile Ser Ile Ser Val Asn Leu Leu Leu	
105 110 115	
GGC AGC CAG TCC TCC GGG AGG CCC ACA GTT ACT GCC TCC AGC TGC AGC	480
Gly Ser Glu Ser Ser Gly Arg Pro Thr Val Thr Ala Ser Ser Cys Ser	
120 125 130 135	

AGT GAC ATC GCT GAC GTG GAG GTG GAC ATG TCG GCA GAC TTG GGG TGG Ser Asp Ile Ala Asp Val Glu Val Asp Met Ser Gly Asp Leu Gly Trp	528
140 145 150	
CTG TTG AAC CTC TTC CAC AAC CAG ATT GAG TCC AAG TTC CAG AAA GTA Leu Leu Asn Leu Phe His Asn Gln Ile Glu Ser Lys Phe Gln Lys Val	576
155 160 165	
CTG GAG AGC AGG ATT TGC GAA ATG ATC CAG AAA TCG GTG TCC TCC GAT Leu Glu Ser Arg Ile Cys Glu Met Ile Gln Lys Ser Val Ser Ser Asp	624
170 175 180	
CTA CAG CCT TAT CTC CAA ACT CTG CCA GTT ACA ACA GAG ATT GAC AGT Leu Gln Pro Tyr Leu Gln Thr Leu Pro Val Thr Thr Glu Ile Asp Ser	672
185 190 195	
TTC GCC GAC ATT GAT TAT AGC TTA GTG GAA GCC CCT CGG GCA ACA GCC Phe Ala Asp Ile Asp Tyr Ser Leu Val Glu Ala Pro Arg Ala Thr Ala	720
200 205 210 215	
CAG ATG CTG GAG GTG ATG TTT AAG GGT GAA ATC TTT CAT CGT AAC CAC Gln Met Leu Glu Val Met Phe Lys Gly Glu Ile Phe His Arg Asn His	768
220 225 230	
CGT TCT CCA GTT ACC CTC CTT GCT GCA GTC ATG AGC CTT CCT GAG GAA Arg Ser Pro Val Thr Leu Leu Ala Ala Val Met Ser Leu Pro Glu Glu	816
235 240 245	
CAC AAC AAA ATG GTC TAC TTT GCC ATC TCG GAT TAT GTC TTC AAC ACG His Asn Lys Met Val Tyr Phe Ala Ile Ser Asp Tyr Val Phe Asn Thr	864
250 255 260	
GCC AGC CTG GTT TAT CAT GAG GAA GGA TAT CTG AAC TTC TCC ATC ACA Ala Ser Leu Val Tyr His Glu Glu Gly Tyr Leu Asn Phe Ser Ile Thr	912
265 270 275	
GAT GAG ATC ATA CCG CCT GAC TCT AAT ATC CCA CTG ACC ACC AAG TCC Asp Glu Met Ile Pro Pro Asp Ser Asn Ile Arg Leu Thr Thr Lys Ser	960
280 285 290 295	
TTC CGA CCC TTC GTC CCA CGG TTA GCC AGG CTC TAC CCC AAC ATG AAC Phe Arg Pro Phe Val Pro Arg Leu Ala Arg Leu Tyr Pro Asn Met Asn	1008
300 305 310	
CTG GAA CTC CAG GGA TCA GTG CCC TCT GCT CCG CTC CTG AAC TTC AGC Leu Glu Leu Gln Gly Ser Val Pro Ser Ala Pro Leu Leu Asn Phe Ser	1056
315 320 325	
CCT GCG AAT CTG TCT GTG GAC CCC TAT ATG GAG ATA GAT GCC TTT GTG Pro Gly Asn Leu Ser Val Asp Pro Tyr Met Glu Ile Asp Ala Phe Val	1104
330 335 340	
CTC CTG CCC AGC TCC AGC AAG GAG CCT GTC TTC CGG CTC AGT GTG GCC Leu Leu Pro Ser Ser Ser Lys Glu Pro Val Phe Arg Leu Ser Val Ala	1152
345 350 355	
ACT AAT GTG TCC GCC ACC TTG ACC TTC AAT ACC AGC AAG ATC ACT GGG Thr Asn Val Ser Ala Thr Leu Thr Phe Asn Thr Ser Lys Ile Thr Gly	1200
360 365 370 375	
TTC CTG AAG CCA GGA AAG GTA AAA GTG GAA CTG AAA GAA TCC AAA GTT Phe Leu Lys Pro Gly Lys Val Lys Val Glu Leu Lys Glu Ser Lys Val	1248
380 385 390	

174

GGA CTA TTC AAT GCA GAG CTG TTG GAA GCG CTC CTC AAC TAT TAC ATC 1296
 Gly Leu Phe Asn Ala Glu Leu Leu Glu Ala Leu Leu Asn Tyr Tyr Ile
 395 400 405

CTT AAC ACC TTC TAC CCC AAG TTC AAT GAT AAG TTG CCC GAA GGC TTC 1344
 Leu Asn Thr Phe Tyr Pro Lys Phe Asn Asp Lys Leu Ala Glu Gly Phe
 410 415 420

CCC CTT CCT CTG CTG AAG CGT GTT CAG CTC TAC GAC CTT GCG CTG CAG 1392
 Pro Leu Pro Leu Leu Lys Arg Val Gln Leu Tyr Asp Leu Gly Leu Gln
 425 430 435

ATC CAT AAG GAC TTC CTG TTC TTG GGT GCC AAT GTC CAA TAC ATG AGA 1440
 Ile His Lys Asp Phe Leu Phe Leu Gly Ala Asn Val Gln Tyr Met Arg
 440 445 450 455

GTT 1443
 Val

(2) INFORMATION FOR SEQ ID NO:98:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 481 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "rLBP"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:98:

Met Gly Ala Leu Ala Arg Ala Leu Pro Ser Ile Leu Leu Ala Leu Leu
 -25 -20 -15 -10

Leu Thr Ser Thr Pro Glu Ala Leu Gly Ala Asn Pro Gly Leu Val Ala
 -5 1 5

Arg Ile Thr Asp Lys Gly Leu Gln Tyr Ala Ala Gln Glu Gly Leu Leu
 10 15 20

Ala Leu Gln Ser Glu Leu Leu Arg Ile Thr Leu Pro Asp Phe Thr Gly
 25 30 35

Asp Leu Arg Ile Pro His Val Gly Arg Gly Arg Tyr Glu Phe His Ser
 40 45 50 55

Leu Asn Ile His Ser Cys Glu Leu Leu His Ser Ala Leu Arg Pro Val
 60 65 70

Pro Gly Gln Gly Leu Ser Leu Ser Ile Ser Asp Ser Ser Ile Arg Val
 75 80 85

Gln Gly Arg Trp Lys Val Arg Lys Ser Phe Phe Lys Leu Gln Gly Ser
 90 95 100

Phe Asp Val Ser Val Lys Gly Ile Ser Ile Ser Val Asn Leu Leu Leu
 105 110 115

Gly Ser Glu Ser Ser Gly Arg Pro Thr Val Thr Ala Ser Ser Cys Ser
 120 125 130 135

175

Ser Asp Ile Ala Asp Val Glu Val Asp Met Ser Gly Asp Leu Gly Trp
 140 145 150
 Leu Leu Asn Leu Phe His Asn Gln Ile Glu Ser Lys Phe Gln Lys Val
 155 160 165
 Leu Glu Ser Arg Ile Cys Glu Met Ile Gln Lys Ser Val Ser Ser Asp
 170 175 180
 Leu Gln Pro Tyr Leu Gln Thr Leu Pro Val Thr Thr Glu Ile Asp Ser
 185 190 195
 Phe Ala Asp Ile Asp Tyr Ser Leu Val Glu Ala Pro Arg Ala Thr Ala
 200 205 210 215
 Gln Met Leu Glu Val Met Phe Lys Gly Glu Ile Phe His Arg Asn His
 220 225 230
 Arg Ser Pro Val Thr Leu Leu Ala Ala Val Met Ser Leu Pro Glu Glu
 235 240 245
 His Asn Lys Met Val Tyr Phe Ala Ile Ser Asp Tyr Val Phe Asn Thr
 250 255 260
 Ala Ser Leu Val Tyr His Glu Glu Gly Tyr Leu Asn Phe Ser Ile Thr
 265 270 275
 Asp Glu Met Ile Pro Pro Asp Ser Asn Ile Arg Leu Thr Thr Lys Ser
 280 285 290 295
 Phe Arg Pro Phe Val Pro Arg Leu Ala Arg Leu Tyr Pro Asn Met Asn
 300 305 310
 Leu Glu Leu Gln Gly Ser Val Pro Ser Ala Pro Leu Leu Asn Phe Ser
 315 320 325
 Pro Gly Asn Leu Ser Val Asp Pro Tyr Met Glu Ile Asp Ala Phe Val
 330 335 340
 Leu Leu Pro Ser Ser Ser Lys Glu Pro Val Phe Arg Leu Ser Val Ala
 345 350 355
 Thr Asn Val Ser Ala Thr Leu Thr Phe Asn Thr Ser Lys Ile Thr Gly
 360 365 370 375
 Phe Leu Lys Pro Gly Lys Val Lys Val Glu Leu Lys Glu Ser Lys Val
 380 385 390
 Gly Leu Phe Asn Ala Glu Leu Leu Glu Ala Leu Leu Asn Tyr Tyr Ile
 395 400 405
 Leu Asn Thr Phe Tyr Pro Lys Phe Asn Asp Lys Leu Ala Glu Gly Phe
 410 415 420
 Pro Leu Pro Leu Leu Lys Arg Val Gln Leu Tyr Asp Leu Gly Leu Gln
 425 430 435
 Ile His Lys Asp Phe Leu Phe Leu Gly Ala Asn Val Gln Tyr Met Arg
 440 445 450 455
 Val

(2) INFORMATION FOR SEQ ID NO:99:

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.57-

Cys Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Pro Leu
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.75"

Ile Lys Lys Arg Ala Ile Ser Phe Leu Gly Lys Lys Trp Gln Lys
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.282"

Lys Trp Lys Ala Phe Phe Arg Phe Leu Lys Lys Trp Lys Ala Phe
1 5 10 15

(1) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 16 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

177

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.103"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:102:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Trp Lys Arg Phe Leu Lys
1 5 10 15
Lys

(2) INFORMATION FOR SEQ ID NO:103:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.104"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:103:

Lys Ser Lys Val Gly Trp Leu Ile Ser Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:104:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 16 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.105"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 13
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 13 is beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:104:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Trp Lys Arg Ala Leu Lys
1 5 10 15
Lys

(2) INFORMATION FOR SEQ ID NO:105:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

178

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.106"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:105:

Lys Ser Lys Val Gly Trp Leu Ile Thr Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:106:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.107"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:106:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe Trp Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:107:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.108"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:107:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Trp
1 5 10

(2) INFORMATION FOR SEQ ID NO:108:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.109"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 11
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 11 is beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:108:

179

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Ala His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:109:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.110"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 12
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 12 is beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:109:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe Ala Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:110:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.111"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 14
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 14 is beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:110:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Ala
1 5 10

(2) INFORMATION FOR SEQ ID NO:111:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.112"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 7
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 7 is beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 11
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 11 is beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:111:

Ile Lys Ile Ser Gly Lys Ala Lys Ala Gln Ala Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:112:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.113"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:112:

Lys Ser Lys Val Gly Trp Leu Ile Gln Phe Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:113:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.114"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:113:

Lys Trp Gln Leu Arg Ser Lys Gly Lys Ile Lys Ile Phe Lys Ala
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:114:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.116"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 6 is beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:114:

Lys Ser Lys Val Lys Ala Leu Ile Gln Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:115:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.119"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 7
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 7 is beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 10
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 10 is beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:115:

Ile Lys Ile Ser Gly Lys Ala Lys Ala Ala Lys Arg Phe Leu Lys
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:116:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.120"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:116:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Lys Leu Lys
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:117:

182

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.121"

- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 10
 - (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 10 is beta-1-naphthyl-substituted."

- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 11
 - (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 11 is beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:117:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Ala Ala Arg Phe Leu Lys
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:118:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.122"

- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 7
 - (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 7 is beta-1-naphthyl-substituted."

- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 10
 - (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 10 is beta-1-naphthyl-substituted."

- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 11
 - (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 11 is beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:118:

Ile Lys Ile Ser Gly Lys Ala Lys Ala Ala Ala Arg Phe Leu Lys
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO:119:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.123"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 9
- (C) OTHER INFORMATION: /label= Substituted-Phe
 /note= "The phenylalanine at position 9 is
 p-amino-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:119:

Lys Ser Lys Val Gly Trp Leu Ile Phe Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:120:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.124"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:120:

Lys Ser Lys Val Lys Trp Leu Ile Gln Leu Trp His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:121:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.125"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:121:

Lys Ser Lys Val Gly Trp Leu Ile Tyr Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:122:

184

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.126"

- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6
(C) OTHER INFORMATION: /label= D-Trp
/note= "The amino acid at position 6 is
D-tryptophan."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:122:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:123:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.127"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:123:

Lys Ser Lys Val Gly Phe Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:124:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc_feature
(D) OTHER INFORMATION: "BPI.128"

- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6
(C) OTHER INFORMATION: /label= D-Phe
/note= "The amino acid at position 6 is
D-phenylalanine."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:124:

Lys Ser Lys Val Gly Phe Leu Ile Gln Leu Pro His Lys Lys
1 5 10

185

(2) INFORMATION FOR SEQ ID NO:125:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.129"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 6 is
D-1-beta-1-naphthyl-
substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:125:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:126:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.130"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 6 is
2-beta-1-naphthyl-
substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:126:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:127:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.131"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 6 is
D-2-beta-1-naphthyl-
substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:127:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:128:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.132"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 6 is
pyridyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:128:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:129:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.133"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Phe
/note= "The phenylalanine at position 6 is
para-amino-
substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:129:

Lys Ser Lys Val Gly Phe Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:130:

- (i) SEQUENCE CHARACTERISTICS:

187

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.134"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 5
- (C) OTHER INFORMATION: /label= Substituted-Phe
/note= "The phenylalanine at position 5 is
para-amino-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:130:

Lys Ser Lys Val Phe Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:131:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.135"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:131:

Lys Ser Lys Val Gly Lys Leu Ile Gln Leu Pro His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:132:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.136"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:132:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Glu Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:133:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 16 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.137"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:133:

Cys Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
1 5 10 15

Cys

(2) INFORMATION FOR SEQ ID NO:134:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.138"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:134:

Lys Ser Lys Val Lys Phe Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:135:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.139"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:135:

Lys Ser Lys Val Gly Tyr Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:136:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 7 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.140"

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 1

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 1 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 2

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 2 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:136:

Ala Ala Arg Phe Leu Lys Phe
1 5

(2) INFORMATION FOR SEQ ID NO:137:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 15 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.141"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:137:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Trp Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:138:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.142"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:138:

Lys Ser Lys Val Gly Trp Leu Ile Gln Trp Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:139:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.143"

190

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 10
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 10 is
beta-l-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:139:

Lys Ser Lys Val Gly Trp Leu Ile Gln Ala Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:140:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.144"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 6 is
cyclohexyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:140:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:141:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 24 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.145"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:141:

Lys Trp Lys Ala Ala Ala Arg Phe Leu Lys Lys Ser Lys Val Gly
1 5 10 15

Trp Leu Ile Gln Leu Phe His Lys Lys
20

(2) INFORMATION FOR SEQ ID NO:142:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.146"

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 12
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 12 is
beta-1-naphthyl-substituted."

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 14
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 14 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:142:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe Ala Lys Ala
1 5 10

(2) INFORMATION FOR SEQ ID NO:143:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.147"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:143:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Glu Lys Lys Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:144:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.148"

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 6 is
beta-1-naphthyl-substituted."

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 12

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 12 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:144:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe Ala Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:145:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 1813 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE:
(A) NAME/KEY: CDS
(B) LOCATION: 31..1491

(ix) FEATURE:
(A) NAME/KEY: mat_peptide
(B) LOCATION: 124..1491

(ix) FEATURE:
(A) NAME/KEY: misc_feature
(D) OTHER INFORMATION: "rBPI"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:145:

CAGGCCTTGA GGTTTTGGCA GCTCTGGAGG ATG ACA GAG AAC ATG GCC AGG GGC	54
Met Arg Glu Asn Met Ala Arg Gly	
-31 -30 -25	
CCT TGC AAC GCG CCG ACA TCG GTG TCC CTG ATG GTG CTC GTC GCC ATA	102
Pro Cys Asn Ala Pro Arg Trp Val Ser Leu Met Val Leu Val Ala Ile	
-20 -15 -10	
GCC ACC GCC GTG ACA GCG GCC GTC AAC CCT GGC GTC GTG GTC AGG ATC	150
Gly Thr Ala Val Thr Ala Ala Val Asn Pro Gly Val Val Val Arg Ile	
-5 1 5	
TCC CAG AAG GCG CTG GAC TAC GCC AGC CAG CAG GGG ACG GCC GCT CTG	198
Ser Gln Lys Gly Leu Asp Tyr Ala Ser Gln Gln Gly Thr Ala Ala Leu	
10 15 20 25	
CAG AAG GAG CTG AAG AGG ATC AAG ATT CCT GAC TAC TCA GAC AGC TTT	246
Gln Lys Glu Leu Lys Arg Ile Lys Ile Pro Asp Tyr Ser Asp Ser Phe	
30 35 40	
AAG ATC AAG CAT CTT GCG AAG GCG CAT TAT AGC TTC TAC AGC ATG GAC	294
Lys Ile Lys His Leu Gly Lys Gly His Tyr Ser Phe Tyr Ser Met Asp	
45 50 55	
ATC CGT GAA TTC CAG CTT CCC AGT TCC CAG ATA AGC ATG GTG CCC AAT	342
Ile Arg Glu Phe Gln Leu Pro Ser Ser Gln Ile Ser Met Val Pro Asn	
60 65 70	
GTG GCG CTT AAG TTC TCC ATC AGC AAC GCC AAT ATC AAG ATC AGC GCG	390
Val Gly Leu Lys Phe Ser Ile Ser Asn Ala Asn Ile Lys Ile Ser Gly	
75 80 85	

AAA TGG AAG GCA CAA AAG AGA TTC TTA AAA ATG AGC GGC AAT TTT GAC Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys Met Ser Gly Asn Phe Asp 90 95 100 105	438
CTG AGC ATA GAA GGC ATG TCC ATT TCG GCT GAT CTG AAG CTG GGC AGT Leu Ser Ile Glu Gly Met Ser Ile Ser Ala Asp Leu Lys Leu Gly Ser 110 115 120	486
AAC CCC ACC TCA GGC AAG CCC ACC ATC ACC TGC TCC AGC TGC AGC AGC Asn Pro Thr Ser Gly Lys Pro Thr Ile Thr Cys Ser Ser Cys Ser Ser 125 130 135	534
CAC ATC AAC AGT GTC CAC GTG CAC ATC TCA AAG AGC AAA GTC GGG TGG His Ile Asn Ser Val His Val His Ile Ser Lys Ser Lys Val Gly Trp 140 145 150	582
CTG ATC CAA CTC TTC CAC AAA AAA ATT GAG TCT GCG CTT CGA AAC AAG Leu Ile Gln Leu Phe His Lys Lys Ile Glu Ser Ala Leu Arg Asn Lys 155 160 165	630
ATG AAC AGC CAG GTC TGC GAG AAA GTG ACC AAT TCT GTA TCC TCC AAG Met Asn Ser Gln Val Cys Glu Lys Val Thr Asn Ser Val Ser Ser Lys 170 175 180 185	678
CTG CAA CCT TAT TTC CAG ACT CTG CCA GTA ATG ACC AAA ATA GAT TCT Leu Gln Pro Tyr Phe Gln Thr Leu Pro Val Met Thr Lys Ile Asp Ser 190 195 200	726
GTG GCT GGA ATC AAC TAT GGT CTG GTG GCA CCT CCA GCA ACC ACG GCT Val Ala Gly Ile Asn Tyr Gly Leu Val Ala Pro Pro Ala Thr Thr Ala 205 210 215	774
GAG ACC CTG GAT GTA CAG ATG AAG GGG GAG TTT TAC AGT GAG AAC CAC Glu Thr Leu Asp Val Gln Met Lys Gly Glu Phe Tyr Ser Glu Asn His 220 225 230	822
CAC AAT CCA CCT CCC TTT GCT CCA CCA GTG ATG GAG TTT CCC GCT GCC His Asn Pro Pro Pro Phe Ala Pro Pro Val Met Glu Phe Pro Ala Ala 235 240 245	870
CAT GAC CGC ATG GTA TAC CTG GGC CTC TCA GAC TAC TTC TTC AAC ACA His Asp Arg Met Val Tyr Leu Gly Leu Ser Asp Tyr Phe Phe Asn Thr 250 255 260 265	918
CCC GCG CTT GTA TAC CAA GAG GCT GCG GTG TTG AAG ATG ACC CTT AGA Ala Gly Leu Val Tyr Gln Glu Ala Gly Val Leu Lys Met Thr Leu Arg 270 275 280	966
GAT GAC ATG ATT CCA AAG GAG TCC AAA TTT CGA CTG ACA ACC AAG TTC Asp Asp Met Ile Pro Lys Glu Ser Lys Phe Arg Leu Thr Thr Lys Phe 285 290 295	1014
TTT CGA ACC TTC CTA CCT GAG GTG GCC AAG AAG TTT CCC AAC ATG AAG Phe Gly Thr Phe Leu Pro Glu Val Ala Lys Lys Phe Pro Asn Met Lys 300 305 310	1062
ATA CAG ATC CAT GTC TCA GCC TCC ACC CCG CCA CAC CTG TCT GTG CAG Ile Gln Ile His Val Ser Ala Ser Thr Pro Pro His Leu Ser Val Gln 315 320 325	1110
CCC ACC GCG CTT ACC TTC TAC CCT GCC GTG GAT GTC CAG GCC TTT GCC Pro Thr Gly Leu Thr Phe Tyr Pro Ala Val Asp Val Gln Ala Phe Ala 330 335 340 345	1158

GTC CTC CCC AAC TCC TCC CTG GCT TCC CTC TTC CTG ATT GGC ATG CAC Val Leu Pro Asn Ser Ser Leu Ala Ser Leu Phe Leu Ile Gly Met His 350 355 360	1206
ACA ACT GGT TCC ATG GAG GTC AGC GCC GAG TCC AAC AGG CTT GTT GGA Thr Thr Gly Ser Met Glu Val Ser Ala Glu Ser Asn Arg Leu Val Gly 365 370 375	1254
GAG CTC AAG CTG GAT AGG CTG CTC CTG GAA CTG AAG CAC TCA AAT ATT Glu Leu Lys Leu Asp Arg Leu Leu Leu Glu Leu Lys His Ser Asn Ile 380 385 390	1302
GGC CCC TTC CCG GTT CAA TTG CTG CAG GAT ATC ATG AAC TAC ATT GTA Gly Pro Phe Pro Val Glu Leu Leu Gln Asp Ile Met Asn Tyr Ile Val 395 400 405	1350
CCC ATT CTT GTG CTG CCC AGG GTT AAC GAG AAA CTA CAG AAA GGC TTC Pro Ile Leu Val Leu Pro Arg Val Asn Glu Lys Leu Gln Lys Gly Phe 410 415 420 425	1398
CCT CTC CCG ACG CCG GCC AGA GTC CAG CTC TAC AAC GTA GTG CTT CAG Pro Leu Pro Thr Pro Ala Arg Val Gln Leu Tyr Asn Val Val Leu Gln 430 435 440	1446
CCT CAC CAG AAC TTC CTG CTC TTC GGT GCA GAC GTT GTC TAT AAA Pro His Gln Asn Phe Leu Leu Phe Gly Ala Asp Val Val Tyr Lys 445 450 455	1491
TGAAGGCACC ACGGGTGCCC GCGGCTGTCA GCCGCACCTG TTCCTGATGG GCTGTGGGGC	1551
ACCGGCTGCC TTTCCTCAGG GAATCCTCTC CAGATCTTAA CCAAGAGCCC CTTGCAAACT	1611
TCTTCGACTC AGATTACAGAA ATGATCTTAA CACGAGGAAA CATTATTCAT TCGAAAAGTG	1671
CATGGTGTGT ATTTTAGGGA TTATGAGCTT CTTCAAGGG CTAAGGCTGC AGAGATATTT	1731
CCTCCAGGAA TCGTGTTCAT ATTGTAACCA AGAAATTTC ATTGTGCTT CATGAAAAAA	1791
AACITCTGGT TTTTTCATG TG	1813

(2) INFORMATION FOR SEQ ID NO:146:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 487 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:146:

Met Arg Glu Asn Met Ala Arg Gly Pro Cys Asn Ala Pro Arg Trp Val
 -31 -30 -25 -20
 Ser Leu Met Val Leu Val Ala Ile Gly Thr Ala Val Thr Ala Ala Val
 -15 -10 -5 1
 Asn Pro Gly Val Val Val Arg Ile Ser Gln Lys Gly Leu Asp Tyr Ala
 5 10 15
 Ser Gln Gln Gly Thr Ala Ala Leu Gln Lys Glu Leu Lys Arg Ile Lys
 20 25 30
 Ile Pro Asp Tyr Ser Asp Ser Phe Lys Ile Lys His Leu Gly Lys Gly
 35 40 45

195

His Tyr Ser Phe Tyr Ser Met Asp Ile Arg Glu Phe Gln Leu Pro Ser
 50 55 60 65
 Ser Gln Ile Ser Met Val Pro Asn Val Gly Leu Lys Phe Ser Ile Ser
 70 75 80
 Asn Ala Asn Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Phe
 85 90 95
 Leu Lys Met Ser Gly Asn Phe Asp Leu Ser Ile Glu Gly Met Ser Ile
 100 105 110
 Ser Ala Asp Leu Lys Leu Gly Ser Asn Pro Thr Ser Gly Lys Pro Thr
 115 120 125
 Ile Thr Cys Ser Ser Cys Ser Ser His Ile Asn Ser Val His Val His
 130 135 140 145
 Ile Ser Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
 150 155 160
 Ile Glu Ser Ala Leu Arg Asn Lys Met Asn Ser Gln Val Cys Glu Lys
 165 170 175
 Val Thr Asn Ser Val Ser Ser Lys Leu Gln Pro Tyr Phe Gln Thr Leu
 180 185 190
 Pro Val Met Thr Lys Ile Asp Ser Val Ala Gly Ile Asn Tyr Gly Leu
 195 200 205
 Val Ala Pro Pro Ala Thr Thr Ala Glu Thr Leu Asp Val Gln Met Lys
 210 215 220 225
 Gly Glu Phe Tyr Ser Glu Asn His His Asn Pro Pro Phe Ala Pro
 230 235 240
 Pro Val Met Glu Phe Pro Ala Ala His Asp Arg Met Val Tyr Leu Gly
 245 250 255
 Leu Ser Asp Tyr Phe Phe Asn Thr Ala Gly Leu Val Tyr Gln Glu Ala
 260 265 270
 Gly Val Leu Lys Met Thr Leu Arg Asp Asp Met Ile Pro Lys Glu Ser
 275 280 285
 Lys Phe Arg Leu Thr Thr Lys Phe Phe Gly Thr Phe Leu Pro Glu Val
 290 295 300 305
 Ala Lys Lys Phe Pro Asn Met Lys Ile Gln Ile His Val Ser Ala Ser
 310 315 320
 Thr Pro Pro His Leu Ser Val Gln Pro Thr Gly Leu Thr Phe Tyr Pro
 325 330 335
 Ala Val Asp Val Gln Ala Phe Ala Val Leu Pro Asn Ser Ser Leu Ala
 340 345 350
 Ser Leu Phe Leu Ile Gly Met His Thr Thr Gly Ser Met Glu Val Ser
 355 360 365
 Ala Glu Ser Asn Arg Leu Val Gly Glu Leu Lys Leu Asp Arg Leu Leu
 370 375 380 385
 Leu Glu Leu Lys His Ser Asn Ile Gly Pro Phe Pro Val Glu Leu Leu
 390 395 400

Gln Asp Ile Met Asn Tyr Ile Val Pro Ile Leu Val Leu Pro Arg Val
 405 410 415
 Asn Glu Lys Leu Gln Lys Gly Phe Pro Leu Pro Thr Pro Ala Arg Val
 420 425 430
 Gln Leu Tyr Asn Val Val Leu Gln Pro His Gln Asn Phe Leu Leu Phe
 435 440 445
 Gly Ala Asp Val Val Tyr Lys
 450 455

(2) INFORMATION FOR SEQ ID NO:147:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 24 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.149"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:147:

Lys Trp Lys Val Phe Lys Lys Ile Glu Lys Lys Ser Lys Val Gly
 1 5 10 15
 Trp Leu Ile Gln Leu Phe His Lys Lys
 20

(2) INFORMATION FOR SEQ ID NO:148:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 20 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.150"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:148:

Lys Trp Ala Phe Ala Lys Lys Gln Lys Lys Arg Leu Lys Arg Gln
 1 5 10 15
 Trp Leu Lys Lys Phe
 20

(2) INFORMATION FOR SEQ ID NO:149:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 30 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.153"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:149:

Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys Lys Trp Lys Ala Gln
1 5 10 15
Lys Arg Phe Leu Lys Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys
20 25 30

(2) INFORMATION FOR SEQ ID NO:150:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.154"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 5
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 5 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:150:

Lys Trp Lys Ala Ala Ala Arg Phe Leu Lys Lys Trp Lys Ala Gln
1 5 10 15
Lys Arg Phe Leu Lys
20

(2) INFORMATION FOR SEQ ID NO:151:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.155"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 15
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 15 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 16

(C) OTHER INFORMATION: /label= Substituted-Ala
 /note= "Position 16 is
 beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:151:

Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys Lys Trp Lys Ala Ala
 1 5 10 15
 Ala Arg Phe Leu Lys
 20

(2) INFORMATION FOR SEQ ID NO:152:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 20 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.156"

(ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 5
 (C) OTHER INFORMATION: /label= Substituted-Ala
 /note= "Position 5 is
 beta-1-naphthyl-substituted."

(ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 6
 (C) OTHER INFORMATION: /label= Substituted-Ala
 /note= "Position 6 is
 beta-1-naphthyl-substituted."

(ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 15
 (C) OTHER INFORMATION: /label= Substituted-Ala
 /note= "Position 15 is
 beta-1-naphthyl-substituted."

(ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 16
 (C) OTHER INFORMATION: /label= Substituted-Ala
 /note= "Position 16 is
 beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:152:

Lys Trp Lys Ala Ala Ala Arg Phe Leu Lys Lys Trp Lys Ala Ala
 1 5 10 15
 Ala Arg Phe Leu Lys
 20

(2) INFORMATION FOR SEQ ID NO:153:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 30 amino acids
 (B) TYPE: amino acid

199

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.157"

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 5

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 5 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 6

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 15

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 15 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 16

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 16 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 25

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 25 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 26

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 26 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:153:

Lys	Trp	Lys	Ala	Ala	Ala	Arg	Phe	Leu	Lys	Lys	Trp	Lys	Ala	Ala
1				5					10				15	

Ala	Arg	Phe	Leu	Lys	Lys	Trp	Lys	Ala	Ala	Ala	Arg	Phe	Leu	Lys
			20					25					30	

(2) INFORMATION FOR SEQ ID NO:154:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 29 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

200

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.158"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 10
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 10 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 11
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 11 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:154:

```

Ile Lys Ile Ser Gly Lys Trp Lys Ala Ala Ala Arg Phe Leu Lys
1           5           10           15
Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
                20           25

```

(2) INFORMATION FOR SEQ ID NO:155:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 24 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.159"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 2
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 2 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:155:

```

Lys Ala Lys Ala Gln Ala Arg Phe Leu Lys Lys Ser Lys Val Gly
1           5           10           15
Trp Leu Ile Gln Leu Trp His Lys Lys
                20

```

(2) INFORMATION FOR SEQ ID NO:156:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

201

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.160"

(ix) FEATURE:

(A) NAME/KEY: Modified-site
(B) LOCATION: 2
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 2 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site
(B) LOCATION: 6
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site
(B) LOCATION: 12
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 12 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site
(B) LOCATION: 16
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 16 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:156:

Lys	Ala	Lys	Ala	Gln	Ala	Arg	Phe	Leu	Lys	Lys	Ala	Lys	Ala	Gln
1				5					10				15	
Ala Arg Phe Leu Lys														
20														

(2) INFORMATION FOR SEQ ID NO:157:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.161"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:157:

Lys	Ser	Lys	Val	Lys	Ala	Leu	Ile	Gln	Leu	Phe	His	Lys	Lys
1				5				10					

(2) INFORMATION FOR SEQ ID NO:158:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24 amino acids
(B) TYPE: amino acid

202

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc_feature

(D) OTHER INFORMATION: "BPI.162"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:158:

Lys Trp Lys Ala Gln Trp Arg Phe Leu Lys Lys Ser Lys Val Gly
1 5 10 15
Trp Leu Ile Gln Leu Phe His Lys Lys
20

(2) INFORMATION FOR SEQ ID NO:159:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc_feature

(D) OTHER INFORMATION: "BPI.163"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:159:

Lys Trp Lys Ala Gln Trp Arg Phe Leu Lys Lys Trp Lys Ala Gln
1 5 10 15
Trp Arg Phe Leu Lys
20

(2) INFORMATION FOR SEQ ID NO:160:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc_feature

(D) OTHER INFORMATION: "BPI.164"

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 5

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 5 is
beta-l-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 15

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 15 is
beta-l-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:160:

Lys Trp Lys Ala Ala Lys Arg Phe Leu Lys Lys Trp Lys Ala Ala
 1 5 10 15
 Lys Arg Phe Leu Lys
 20

(2) INFORMATION FOR SEQ ID NO:161:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.165"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 2
- (C) OTHER INFORMATION: /label= Substituted-Ala
 /note= "Position 2 is
 beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 12
- (C) OTHER INFORMATION: /label= Substituted-Ala
 /note= "Position 12 is
 beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:161:

Lys Ala Lys Ala Gln Phe Arg Phe Leu Lys Lys Ala Lys Ala Gln
 1 5 10 15
 Phe Arg Phe Leu Lys
 20

(2) INFORMATION FOR SEQ ID NO:162:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.166"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:162:

Lys Ser Lys Val Gly Val Leu Ile Gln Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:163:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 8 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

204

- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.167"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:163:
Lys Trp Lys Ala Gln Lys Arg Phe
1 5
- (2) INFORMATION FOR SEQ ID NO:164:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: circular
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.168"
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:164:
Cys Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys Met Ser Cys
1 5 10
- (2) INFORMATION FOR SEQ ID NO:165:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: circular
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.169"
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:165:
Cys Lys Trp Lys Ala Gln Lys Arg Phe Cys
1 5 10
- (2) INFORMATION FOR SEQ ID NO:166:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.221"
 - (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 13
 - (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 13 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:166:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Gln Lys Arg Ala Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:167:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.222"

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 14
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 14 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:167:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe His Lys Ala
1 5 10

(2) INFORMATION FOR SEQ ID NO:168:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.223"

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 10
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 10 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:168:

206

Lys Ser Lys Val Gly Ala Leu Ile Gln Ala Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:169:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.224"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 9
- (C) OTHER INFORMATION: /label= Substituted-Phe
/note= "Position 9 is
para-amino-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:169:

Lys Ser Lys Val Gly Ala Leu Ile Phe Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:170:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.225"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 5
- (C) OTHER INFORMATION: /label= Substituted-Phe
/note= "Position 5 is
para-amino-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:170:

Lys Ser Lys Val Phe Ala Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:171:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.226"

- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 6
 - (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:171:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Trp His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:172:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.227"

- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 10
 - (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 10 is
beta-1-naphthyl-substituted."

- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 14
 - (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 14 is
beta-1-naphthyl-substituted."

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:172:

Lys Ser Lys Val Gly Trp Leu Ile Gln Ala Phe His Lys Ala
1 5 10

(2) INFORMATION FOR SEQ ID NO:173:

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- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.228"
- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 9
(C) OTHER INFORMATION: /label= Substituted-Phe
/note= "Position 9 is
para-amino-substituted."
- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 14
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 14 is
beta-1-naphthyl-substituted."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:173:
Lys Ser Lys Val Gly Trp Leu Ile Phe Leu Phe His Lys Ala
1 5 10
- (2) INFORMATION FOR SEQ ID NO:174:
- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.229"
- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 5
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 5 is
para-amino-substituted."
- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 14
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 14 is
beta-1-naphthyl-substituted."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:174:
Lys Ser Lys Val Phe Trp Leu Ile Gln Leu Phe His Lys Ala
1 5 10

(2) INFORMATION FOR SEQ ID NO:175:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.230"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 14
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 14 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:175:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Trp His Lys Ala
1 5 10

(2) INFORMATION FOR SEQ ID NO:176:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.231"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 10
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 10 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 12
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 12 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:176:

Lys Ser Lys Val Gly Trp Leu Ile Gln Ala Phe Ala Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:177:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.232"

(ix) FEATURE:

(A) NAME/KEY: Modified-site
(B) LOCATION: 9
(C) OTHER INFORMATION: /label= Substituted-Phe
/note= "Position 9 is
para-amino-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site
(B) LOCATION: 12
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 12 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:177:

Lys Ser Lys Val Gly Trp Leu Ile Phe Leu Phe Ala Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:178:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.233"

(ix) FEATURE:

(A) NAME/KEY: Modified-site
(B) LOCATION: 5
(C) OTHER INFORMATION: /label= Substituted-Phe
/note= "Position 5 is
para-amino-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site
(B) LOCATION: 12
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 12 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:178:

Lys Ser Lys Val Phe Trp Leu Ile Gln Leu Phe Ala Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:179:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.234"

- (ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 12
 (C) OTHER INFORMATION: /label= Substituted-Ala
 /note= "Position 12 is
 beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:179:

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Trp Ala Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:180:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 14 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.235"

- (ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 9
 (C) OTHER INFORMATION: /label= Substituted-Phe
 /note= "Position 9 is
 para-amino-substituted."

- (ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 10
 (C) OTHER INFORMATION: /label= Substituted-Ala
 /note= "Position 10 is
 beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:180:

Lys Ser Lys Val Gly Trp Leu Ile Phe Ala Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:181:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 14 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.236"

- (ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 5
 (C) OTHER INFORMATION: /label= Substituted-Phe
 /note= "Position 5 is
 para-amino-substituted."

- (ix) FEATURE:
 (A) NAME/KEY: Modified-site

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(B) LOCATION: 10
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 10 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:181:

Lys Ser Lys Val Phe Trp Leu Ile Gln Ala Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:182:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.237"

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 10
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 10 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:182:

Lys Ser Lys Val Gly Trp Leu Ile Gln Ala Trp His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:183:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.238"

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 5
(C) OTHER INFORMATION: /label= Substituted-Phe
/note= "Position 5 is
para-amino-substituted."

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 9
(C) OTHER INFORMATION: /label= Substituted-Phe
/note= "Position 9 is
para-amino-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:183:

Lys Ser Lys Val Phe Trp Leu Ile Phe Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:184:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.239"

- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 9
(C) OTHER INFORMATION: /label= Substituted-Phe
/note= "Position 9 is
para-amino-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:184:

Lys Ser Lys Val Gly Trp Leu Ile Phe Leu Trp His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:185:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.240"

- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 5
(C) OTHER INFORMATION: /label= Substituted-Phe
/note= "Position 5 is
para-amino-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:185:

Lys Ser Lys Val Phe Trp Leu Ile Gln Leu Trp His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:186:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 24 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.247"

- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 2

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 2 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 6

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:186:

Lys Ala Lys Ala Gln Ala Arg Phe Leu Lys Lys Ser Lys Val Gly
1 5 10 15
Trp Leu Ile Leu Leu Phe His Lys Lys
20

(2) INFORMATION FOR SEQ ID NO:187:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.245"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:187:

Lys Trp Lys Ala Gln Phe Arg Phe Leu Lys Lys Ser Lys Val Gly
1 5 10 15
Trp Leu Ile Gln Leu Trp His Lys Lys
20

(2) INFORMATION FOR SEQ ID NO:188:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.246"

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 16

(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 16 is
D-beta-2-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:188:

Lys Trp Lys Ala Gln Phe Arg Phe Leu Lys Lys Ser Lys Val Gly
1 5 10 15

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Ala Leu Ile Gln Leu Phe His Lys Lys
20

(2) INFORMATION FOR SEQ ID NO:189:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 24 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.248"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 2
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 2 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
beta-1-naphthyl-substituted."

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 16
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 16 is
D-beta-2-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:189:

Lys Ala Lys Ala Gln Ala Arg Phe Leu Lys Lys Ser Lys Val Gly
1 5 10 15
Ala Leu Ile Gln Leu Phe His Lys Lys
20

(2) INFORMATION FOR SEQ ID NO:190:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.242"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
D-beta-2-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:190:

216

Lys Ser Lys Val Gly Ala Leu Ile Leu Leu Phe His Lys Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO:191:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 28 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.272"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:191:

Lys Ser Lys Val Gly Trp Leu Ile Leu Leu Phe His Lys Lys Lys
 1 5 10 15
 Ser Lys Val Gly Trp Leu Ile Leu Leu Phe His Lys Lys
 20 25

(2) INFORMATION FOR SEQ ID NO:192:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 28 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.275"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:192:

Lys Ser Lys Val Gly Trp Leu Ile Phe Leu Phe His Lys Lys Lys
 1 5 10 15
 Ser Lys Val Gly Trp Leu Ile Phe Leu Phe His Lys Lys
 20 25

(2) INFORMATION FOR SEQ ID NO:193:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 28 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.270"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:193:

Lys Ser Lys Val Gly Trp Leu Ile Leu Leu Phe His Lys Lys Lys
 1 5 10 15
 Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
 20 25

(2) INFORMATION FOR SEQ ID NO:194:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.271"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:194:

```

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys Lys
1           5           10           15
Ser Lys Val Gly Trp Leu Ile Leu Leu Phe His Lys Lys
                20           25

```

(2) INFORMATION FOR SEQ ID NO:195:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.273"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:195:

```

Lys Ser Lys Val Gly Trp Leu Ile Phe Leu Phe His Lys Lys Lys
1           5           10           15
Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys
                20           25

```

(2) INFORMATION FOR SEQ ID NO:196:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (D) OTHER INFORMATION: "BPI.274"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:196:

```

Lys Ser Lys Val Gly Trp Leu Ile Gln Leu Phe His Lys Lys Lys
1           5           10           15
Ser Lys Val Gly Trp Leu Ile Phe Leu Phe His Lys Lys
                20           25

```

(2) INFORMATION FOR SEQ ID NO:197:

- (i) SEQUENCE CHARACTERISTICS:

(ii) MOLECULE TYPE: peptide

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BFI.276"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:197:

Lys Trp Lys Ala Gln Phe Arg Phe Leu Lys Lys Ser Lys Val Gly
1 5 10 15

Trp Leu Ile Phe Leu Phe His Lys Lys
20

(2) INFORMATION FOR SEQ ID NO:198:

(A) LENGTH: 14 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.241"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:198:

Lys Ser Lys Val Gly Trp Leu Ile Leu Leu Trp His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:199:

(A) LENGTH: 14 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.243"

(A) NAME/KEY: Modified-site

(B) LOCATION: 6

(C) OTHER INFORMATION: /label= Substituted-Ala

```
/note= "Position 6 is
```

D-beta-2-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:199:

Lys Ser Lys Val Gly Ala Leu Ile Cln Leu Trp His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:200:

(A) LENGTH: 14 amino acids

(B) TYPE: amino acid

- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "SPI.244"
- (ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6
(C) OTHER INFORMATION: /label= Substituted-Ala
/note= "Position 6 is
D-beta-2-naphthyl-substituted."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:200:
Lys Ser Lys Val Gly Ala Leu Ile Leu Leu Trp His Lys Lys
1 5 10
- (2) INFORMATION FOR SEQ ID NO:201:
- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "SPI.249"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:201:
Lys Ser Lys Val Gly Gly Leu Ile Gln Leu Phe His Lys Lys
1 5 10
- (2) INFORMATION FOR SEQ ID NO:202:
- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "SPI.250"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:202:
Lys Ser Lys Val Gly Leu Leu Ile Gln Leu Phe His Lys Lys
1 5 10
- (2) INFORMATION FOR SEQ ID NO:203:
- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:

220

- (A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.251"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:203:

Lys Ser Lys Val Gly Ile Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:204:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.252"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
(B) LOCATION: 6
(D) OTHER INFORMATION: /label= D-Ala
/note= "The amino acid at position 6 is
D-alanine"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:204:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:205:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.253"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
(B) LOCATION: 6
(D) OTHER INFORMATION: /label= D-Val
/note= "The amino acid at position 6 is
D-valine"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:205:

Lys Ser Lys Val Gly Val Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:206:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

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- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.254"
- (ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 6
 (D) OTHER INFORMATION: /label= beta-Ala
 /note= "The amino acid at position 6 is
 beta-alanine"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:206:
 Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe His Lys Lys
 1 5 10
- (2) INFORMATION FOR SEQ ID NO:207:
- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 14 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.255"
- (ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 6
 (D) OTHER INFORMATION: /label= delta-aba
 /note= "The amino acid at position 6 is
 delta-aminobutyric acid"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:207:
 Lys Ser Lys Val Gly Xaa Leu Ile Gln Leu Phe His Lys Lys
 1 5 10
- (2) INFORMATION FOR SEQ ID NO:208:
- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 14 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 (A) NAME/KEY: misc feature
 (D) OTHER INFORMATION: "BPI.256"
- (ix) FEATURE:
 (A) NAME/KEY: Modified-site
 (B) LOCATION: 6
 (D) OTHER INFORMATION: /label= gaba
 /note= "The amino acid at position 6 is
 gamma-aminobutyric acid"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:208:
 Lys Ser Lys Val Gly Xaa Leu Ile Gln Leu Phe His Lys Lys
 1 5 10

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(2) INFORMATION FOR SEQ ID NO:209:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "SPI.257"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (D) OTHER INFORMATION: /label= d-methyl-A
/note= "The amino acid at position 6 is
delta-Methyl-alanine"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:209:

Lys Ser Lys Val Gly Ala Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:210:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "SPI.258"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6
- (D) OTHER INFORMATION: /label= t-butyl-G
/note= "The amino acid at position 6 is
tert-butyl-glycine"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:210:

Lys Ser Lys Val Gly Gly Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:211:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "SPI.259"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 6

223

(D) OTHER INFORMATION: /label= N-methyl-G
/note= "The amino acid at position 6 is
N-Methyl-glycine"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:211:

Lys Ser Lys Val Gly Gly Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:212:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.260"

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6
(D) OTHER INFORMATION: /label= N-methyl-V
/note= "The amino acid at position 6 is
N-Methyl-valine"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:212:

Lys Ser Lys Val Gly Val Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:213:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.261"

(ix) FEATURE:
(A) NAME/KEY: Modified-site
(B) LOCATION: 6
(D) OTHER INFORMATION: /label= N-methyl-L
/note= "The amino acid at position 6 is
N-Methyl-leucine"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:213:

Lys Ser Lys Val Gly Leu Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:214:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

224

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.262"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:214:

Lys Ser Lys Val Gly Trp Leu Ile Asn Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:215:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.263"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:215:

Lys Ser Lys Val Gly Trp Leu Ile Glu Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:216:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.264"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:216:

Lys Ser Lys Val Gly Trp Leu Ile Asp Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:217:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: misc feature

(D) OTHER INFORMATION: "BPI.265"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:217:

Lys Ser Lys Val Gly Trp Leu Ile Lys Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:218:

225

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.266"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:218:

Lys Ser Lys Val Lys Val Leu Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:219:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.267"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:219:

Lys Ser Lys Val Lys Trp Ala Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:220:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.268"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:220:

Lys Ser Lys Val Gly Val Ala Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:221:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
(A) NAME/KEY: misc feature
(D) OTHER INFORMATION: "BPI.269"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:221:

226

Lys Ser Lys Val Lys Val Ala Ile Gln Leu Phe His Lys Lys
1 5 10

(2) INFORMATION FOR SEQ ID NO:222:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 24 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc_feature
- (D) OTHER INFORMATION: "BPI.277"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 2
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 2 is
beta-1-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:222:

Lys Ala Lys Ala Gln Phe Arg Phe Leu Lys Lys Ser Lys Val Gly
1 5 10 15
Trp Leu Ile Leu Leu Phe His Lys Lys
20

(2) INFORMATION FOR SEQ ID NO:223:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc_feature
- (D) OTHER INFORMATION: "BPI.278"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:223:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Ala Trp Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:224:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc_feature
- (D) OTHER INFORMATION: "BPI.279"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 10
- (C) OTHER INFORMATION: /label= Substituted-Ala

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/note= "The alanine at position 10 is
beta-l-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:224:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Ala Phe Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:225:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.280"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:225:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Ala Phe Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:226:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.281"

(ix) FEATURE:

- (A) NAME/KEY: Modified-site
- (B) LOCATION: 10
- (C) OTHER INFORMATION: /label= Substituted-Ala
/note= "The alanine at position 10 is
beta-l-naphthyl-substituted."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:226:

Ile Lys Ile Ser Gly Lys Trp Lys Ala Ala Arg Phe Leu Lys
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:227:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 12 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (D) OTHER INFORMATION: "BPI.170"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:227:

Lys Trp Lys Ala Gln Lys Arg Phe Leu Lys Met Ser
1 5 10

WHAT IS CLAIMED IS:

1. Use of a BPI protein product for the manufacture of a medicament for the co-treatment of a gram-negative bacterial infection in mammals with an antibiotic other than a penicillin, cephalosporin, rifampicin, or actinomycin D.
5
2. A use according to claim 1 wherein the medicament is for administration of BPI protein product in a dose of greater than 1 mg/kg and up to 100 mg/kg.
- 10 3. A use according to claim 1 wherein the medicament is for the co-treatment of a gram-negative bacterial infection in mammals with an aminoglycoside antibiotic.
4. A use according to claim 1 wherein the medicament is for
15 improving the therapeutic effectiveness in a gram-negative bacterial infection in mammals of an aminoglycoside antibiotic.
5. A use according to claim 4 wherein the medicament is for increasing the susceptibility to an aminoglycoside antibiotic of gram-negative bacteria
20 involved in the gram-negative bacterial infection.
6. A use according to claim 4 wherein the medicament is for reversing resistance to an aminoglycoside antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.
25
7. A use according to any one of claims 3 to 6 wherein the aminoglycoside antibiotic is selected from the group consisting of amikacin, gentamicin and tobramycin.
- 30 8. A use according to any one of claims 3 or 4 wherein the causative agent of the gram-negative bacterial infection is a bacterial species selected from the group consisting of *Citrobacter*, *Edwardsiella*, *Enterobacter*, *Escherichia*,

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Klebsiella, Morganella, Providencia, Proteus, Pseudomonas, Xanthomonas, Salmonella and *Serratia* species.

9. A use according to claim 1 wherein the medicament is for the
5 co-treatment of a gram-negative bacterial infection in mammals with a sulfonamide or trimethoprim antibiotic.

10. A use according to claim 1 wherein the medicament is for
improving the therapeutic effectiveness in a gram-negative bacterial infection in
10 mammals of a sulfonamide or trimethoprim antibiotic.

11. A use according to claim 10 wherein the medicament is for
increasing the susceptibility to a sulfonamide or trimethoprim antibiotic of gram-
negative bacteria involved in the gram-negative bacterial infection.

12. A use according to claim 10 wherein the medicament is for
reversing resistance to a sulfonamide or trimethoprim antibiotic of gram-negative
bacteria involved in the gram-negative bacterial infection.

13. A use according to any one of claims 9 to 12 wherein the
sulfonamide or trimethoprim antibiotic is the combination antibiotic
trimethoprim/sulfamethoxazole.

14. A use according to any one of claims 9 or 10 wherein the
25 causative agent of the gram-negative bacterial infection is a bacterial species selected
from the group consisting of *Pseudomonas, Xanthomonas, Klebsiella, Enterobacter*
and *Proteus* species.

15. A use according to claim 1 wherein the medicament is for the
30 co-treatment of a gram-negative bacterial infection in mammals with a fluoroquinolone
or quinolone antibiotic.

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16. A use according to claim 1 wherein the medicament is for improving the therapeutic effectiveness in a gram-negative bacterial infection in mammals of a fluoroquinolone or quinolone antibiotic.

5 17. A use according to claim 16 wherein the medicament is for increasing the susceptibility to a fluoroquinolone or quinolone antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

10 18. A use according to claim 16 wherein the medicament is for reversing resistance to a fluoroquinolone or quinolone antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

15 19. A use according to any one of claims 15 to 18 wherein the fluoroquinolone antibiotic is selected from the group consisting of ciprofloxacin, norfloxacin and ofloxacin.

20 20. A use according to any one of claims 15 or 16 wherein the causative agent of the gram-negative bacterial infection is a bacterial species selected from the group consisting of *Pseudomonas*, *Xanthomonas*, *Klebsiella* and *Enterobacter* species.

25 21. A use according to claim 1 wherein the medicament is for the co-treatment of a gram-negative bacterial infection in mammals with a β -lactam antibiotic other than a penicillin or cephalosporin.

22. A use according to claim 1 wherein the medicament is for improving the therapeutic effectiveness in a gram-negative bacterial infection in mammals of a β -lactam antibiotic other than a penicillin or cephalosporin.

30 23. A use according to claim 22 wherein the medicament is for increasing the susceptibility, to a β -lactam antibiotic other than a penicillin or

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cephalosporin, of gram-negative bacteria involved in the gram-negative bacterial infection.

24. A use according to claim 22 wherein the medicament is for
5 reversing resistance, to a β -lactam antibiotic other than a penicillin or cephalosporin, of gram-negative bacteria involved in the gram-negative bacterial infection.

25. A use according to any one of claims 21 to 24 wherein the β -
lactam antibiotic is selected from the group consisting of aztreonam and imipenem.
10

26. A use according to any one of claims 21 or 22 wherein the
causative agent of the gram-negative bacterial infection is a bacterial species selected
from the group consisting of *Pseudomonas*, *Citrobacter*, *Enterobacter*, *Proteus* and
Providencia species.
15

27. A use according to claim 1 wherein the medicament is for the
co-treatment of a gram-negative bacterial infection in mammals with a polymyxin
antibiotic.

28. A use according to claim 1 wherein the medicament is for
20 improving the therapeutic effectiveness in a gram-negative bacterial infection in
mammals of a polymyxin antibiotic.

29. A use according to claim 28 wherein the medicament is for
25 increasing the susceptibility to a polymyxin antibiotic of gram-negative bacteria
involved in the gram-negative bacterial infection.

30. A use according to claim 28 wherein the medicament is for
reversing resistance to a polymyxin antibiotic of gram-negative bacteria involved in
30 the gram-negative bacterial infection.

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31. A use according to any one of claims 27 to 30 wherein the polymyxin antibiotic is polymyxin B.

32. A use according to any one of claims 27 or 28 wherein the
5 causative agent of the gram-negative bacterial infection is a bacterial species selected from the group consisting of *Pseudomonas*, *Escherichia*, *Enterobacter*, and *Providencia* species.

33. In a method for treating gram-negative bacterial infection with
10 an aminoglycoside antibiotic, the improvement comprising the step of concurrently administering BPI protein product in an amount effective to improve the therapeutic effectiveness of the aminoglycoside antibiotic.

34. The method of claim 33 wherein the BPI protein product is
15 administered in an amount effective to increase susceptibility to the aminoglycoside antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

35. The method of claim 33 wherein the BPI protein product is
20 administered in an amount effective to reverse resistance to the aminoglycoside antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

36. The method of claim 33 wherein the aminoglycoside antibiotic is selected from the group consisting of amikacin, gentamicin and tobramycin.

25 37. The method of claim 33 wherein the causative agent of the gram-negative bacterial infection is a bacterial species selected from the group consisting of *Citrobacter*, *Edwardsiella*, *Enterobacter*, *Escherichia*, *Klebsiella*, *Morganella*, *Providencia*, *Proteus*, *Pseudomonas*, *Xanthamonas*, *Salmonella* and *Serratia* species.

30

38. A pharmaceutical composition for treatment of gram-negative bacterial infection comprising an aminoglycoside antibiotic and a BPI protein product

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in an amount effective to improve the therapeutic effectiveness of the amin glycoside antibiotic.

39. In a method for treating gram-negative bacterial infection with a sulfonamide or trimethoprim antibiotic, the improvement comprising the step of concurrently administering BPI protein product in an amount effective to improve the therapeutic effectiveness of the sulfonamide or trimethoprim antibiotic.

40. The method of claim 39 wherein the BPI protein product is administered in an amount effective to increase susceptibility to the sulfonamide or trimethoprim antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

41. The method of claim 39 wherein the BPI protein product is administered in an amount effective to reverse resistance to the sulfonamide or trimethoprim antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

42. The method of claim 39 wherein the sulfonamide or trimethoprim antibiotic is the combination antibiotic trimethoprim/sulfamethoxazole.

43. The method of claim 39 wherein the causative agent of the gram-negative bacterial infection is a bacterial species selected from the group consisting of *Pseudomonas*, *Xanthamonas*, *Klebsiella*, *Enterobacter* and *Proteus* species.

44. A pharmaceutical composition for treatment of gram-negative bacterial infection comprising a sulfonamide or trimethoprim antibiotic and a BPI protein product in an amount effective to improve the therapeutic effectiveness of the sulfonamide or trimethoprim antibiotic.

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45. In a method for treating gram-negative bacterial infection with a fluoroquinolone or quinolone antibiotic, the improvement comprising the step of concurrently administering BPI protein product in an amount effective to improve the therapeutic effectiveness of the fluoroquinolone or quinolone antibiotic.

5

46. The method of claim 45 wherein the BPI protein product is administered in an amount effective to increase susceptibility to the fluoroquinolone or quinolone antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

10

47. The method of claim 45 wherein the BPI protein product is administered in an amount effective to reverse resistance to the fluoroquinolone or quinolone antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

15

48. The method of claim 45 wherein the fluoroquinolone antibiotic is selected from the group consisting of ciprofloxacin, norfloxacin and ofloxacin.

49. The method of claim 45 wherein the causative agent of the gram-negative bacterial infection is a bacterial species selected from the group consisting of *Pseudomonas*, *Xanthomonas*, *Klebsiella* and *Enterobacter* species.

20

50. A pharmaceutical composition for treatment of gram-negative bacterial infection comprising a fluoroquinolone or quinolone antibiotic and a BPI protein product in an amount effective to improve the therapeutic effectiveness of the fluoroquinolone or quinolone antibiotic.

25

51. In a method for treating gram-negative bacterial infection with an aztreonam or imipenem antibiotic, the improvement comprising the step of concurrently administering BPI protein product in an amount effective to improve the therapeutic effectiveness of the aztreonam or imipenem antibiotic.

30

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52. The method of claim 51 wherein the BPI protein product is administered in an amount effective to increase susceptibility to the aztreonam or imipenem antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

5

53. The method of claim 51 wherein the BPI protein product is administered in an amount effective to reverse resistance to the aztreonam or imipenem antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

10

54. The method of claim 51 wherein the causative agent of the gram-negative bacterial infection is a bacterial species selected from the group consisting of *Pseudomonas*, *Citrobacter*, *Enterobacter*, *Proteus* and *Providencia* species.

15

55. A pharmaceutical composition for treatment of gram-negative bacterial infection comprising an aztreonam or imipenem antibiotic and a BPI protein product in an amount effective to improve the therapeutic effectiveness of the aztreonam or imipenem antibiotic.

20

56. In a method for treating gram-negative bacterial infection with a polymyxin antibiotic, the improvement comprising the step of concurrently administering BPI protein product in an amount effective to improve the therapeutic effectiveness of the polymyxin antibiotic.

25

57. The method of claim 56 wherein the BPI protein product is administered in an amount effective to increase susceptibility to the polymyxin antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

30

58. The method of claim 56 wherein the BPI protein product is administered in an amount effective to reverse resistance to the polymyxin antibiotic of gram-negative bacteria involved in the gram-negative bacterial infection.

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59. The method of claim 56 wherein the polymyxin antibiotic is polymyxin B.

60. The method of claim 56 wherein the causative agent of the
5 gram-negative bacterial infection is a bacterial species selected from the group consisting of *Pseudomonas*, *Escherichia*, *Enterobacter*, and *Providencia* species.

61. A pharmaceutical composition for treatment of gram-negative
bacterial infection comprising a polymyxin antibiotic and a BPI protein product in an
10 amount effective to improve the therapeutic effectiveness of the polymyxin antibiotic.

62. In a method for treating gram-negative bacterial infection with a
tetracycline antibiotic, the improvement comprising the step of concurrently
administering BPI protein product in an amount effective to improve the therapeutic
15 effectiveness of the tetracycline antibiotic.

63. A pharmaceutical composition for treatment of gram-negative
bacterial infection comprising a tetracycline antibiotic and a BPI protein product in an
amount effective to improve the therapeutic effectiveness of the tetracycline antibiotic.
20

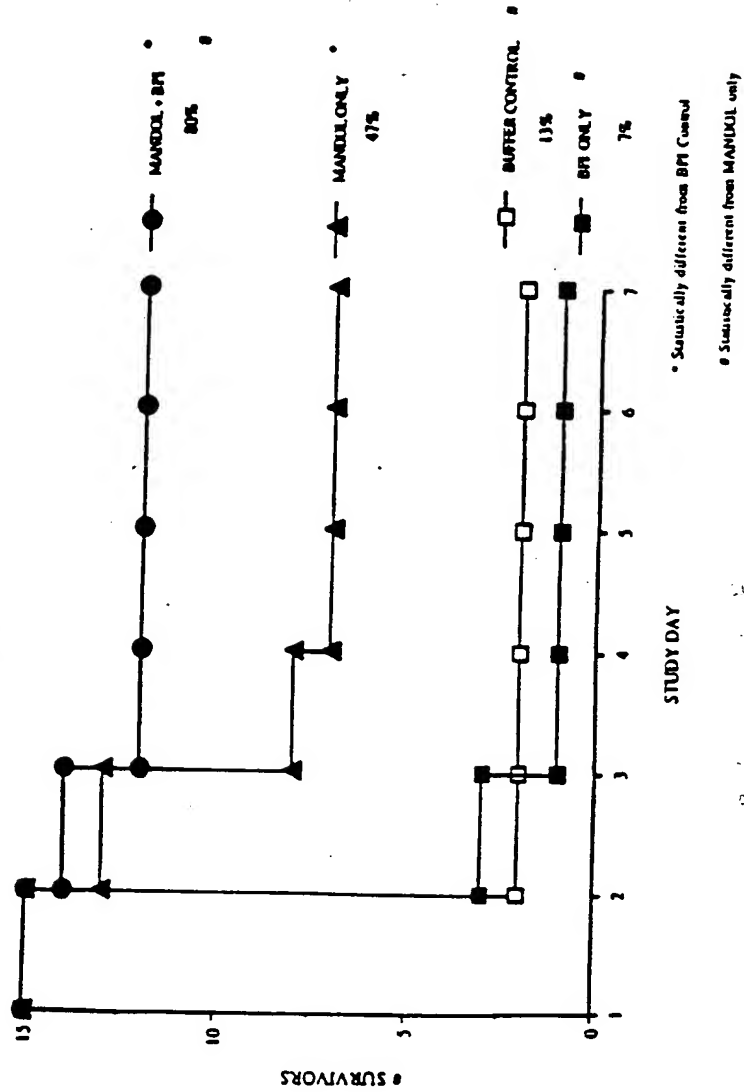
64. A use according to claim 1 wherein the medicament is for the
co-treatment of a gram-negative bacterial infection in mammals with a tetracycline
antibiotic.

25 65. The method of claims 33, 39, 45, 51, 56 or 62 wherein the BPI
protein product is an N-terminal fragment with a molecular weight of approximately
21 to 25 kD, or a dimeric form thereof.

66. The method of claims 33, 39, 45, 51, 56 or 62 wherein the BPI
30 protein product is rBPI₂₃, rBPI₂₁, or rBPI₃₀.

Fig. 1

I.P. E. coli 0111:B4 Bacteria; I.P. MANDOL; and
I.P. BPI delta cys in ICR Mice



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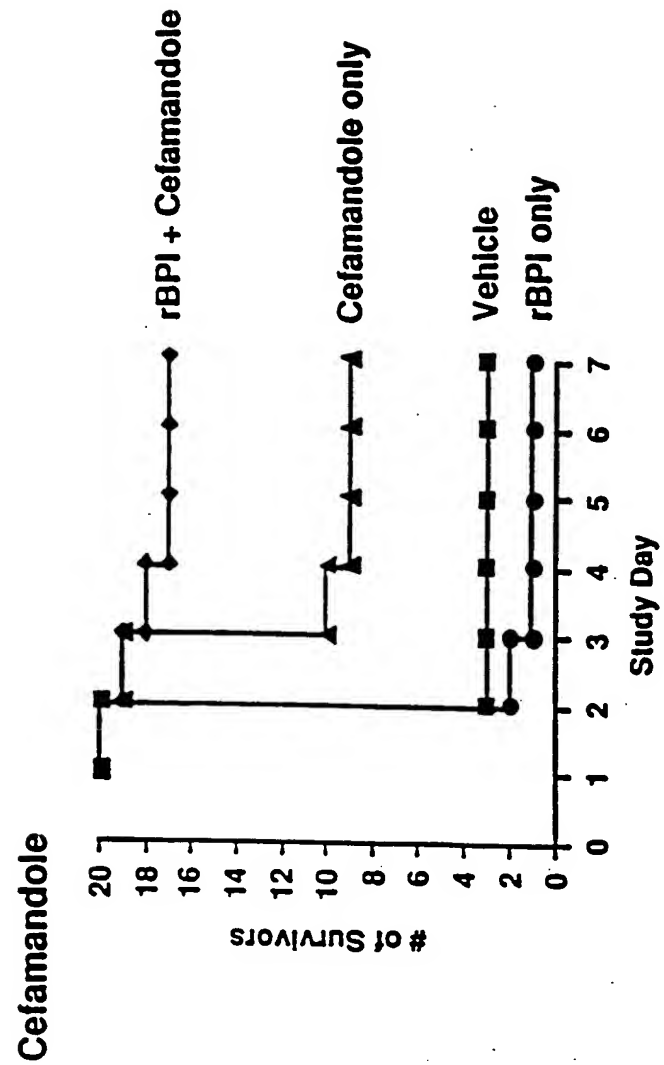


Fig. 2

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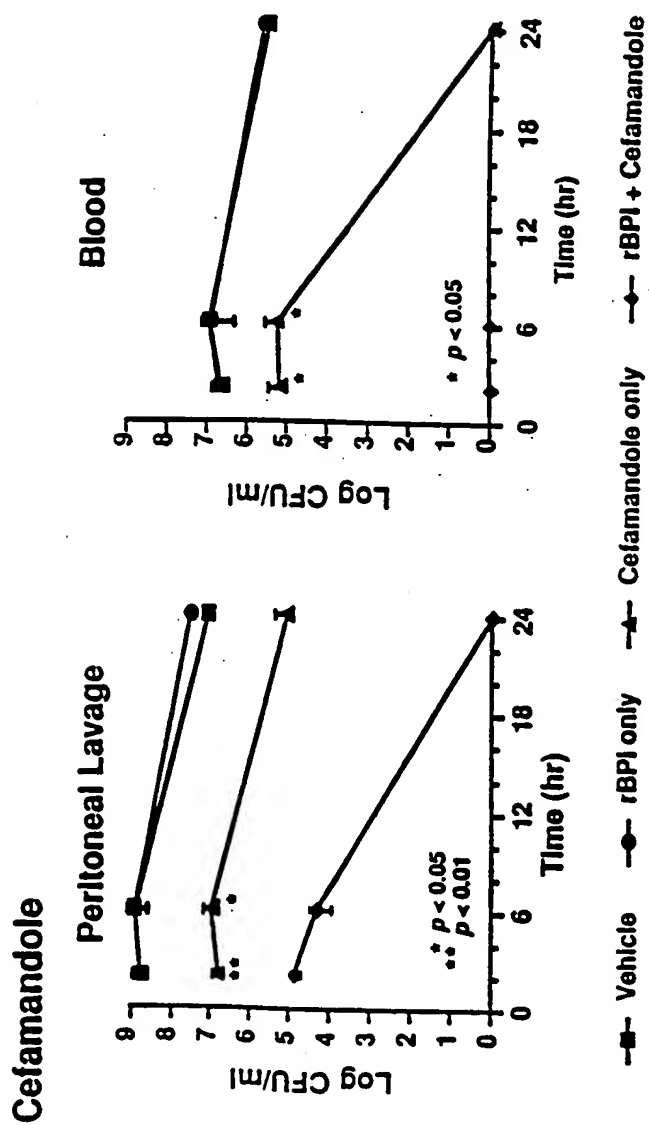


Fig. 3

Fig. 4

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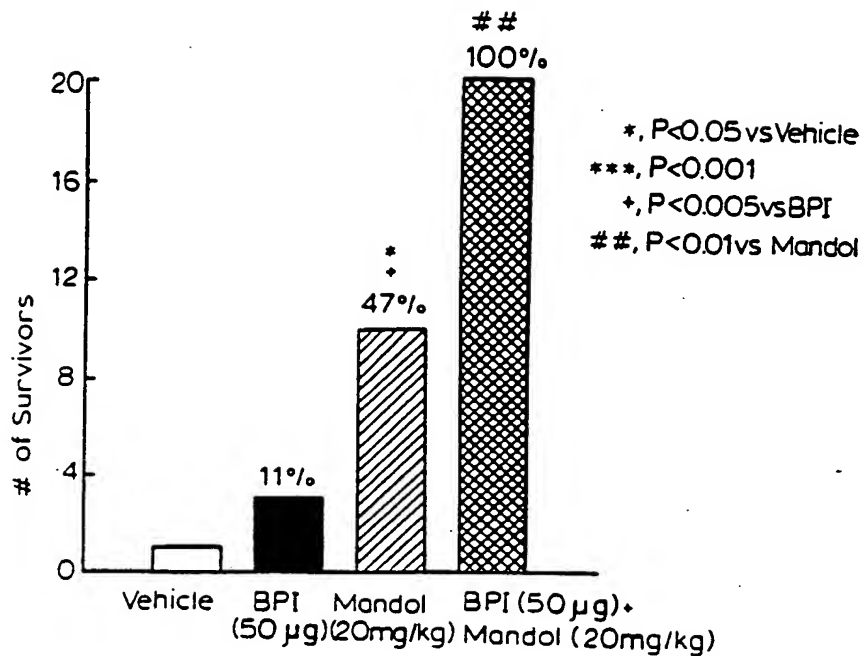


FIG. 5A

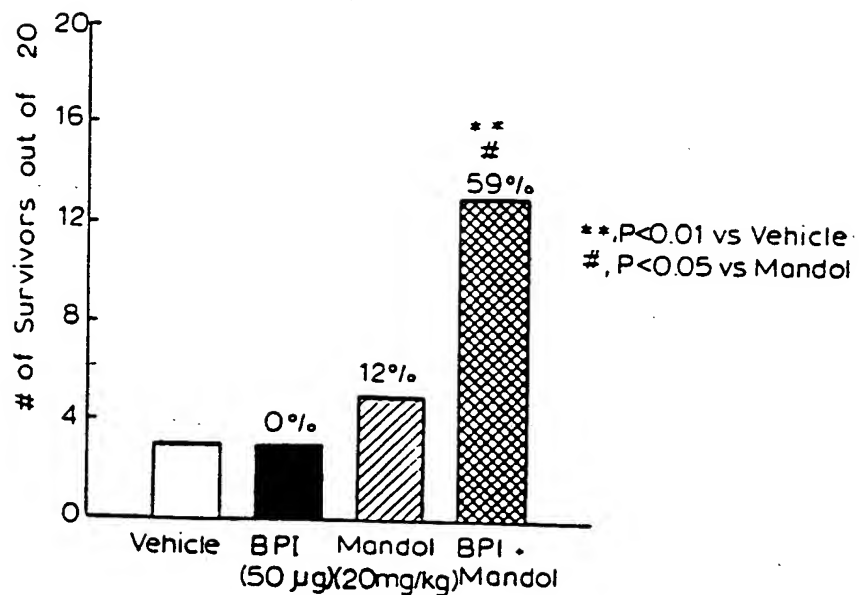


FIG. 5B

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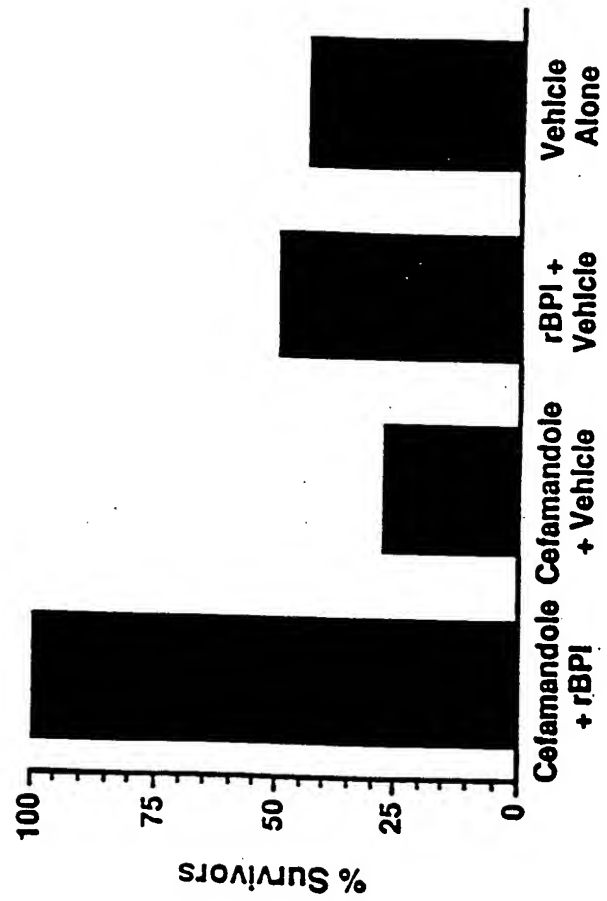


Fig. 6

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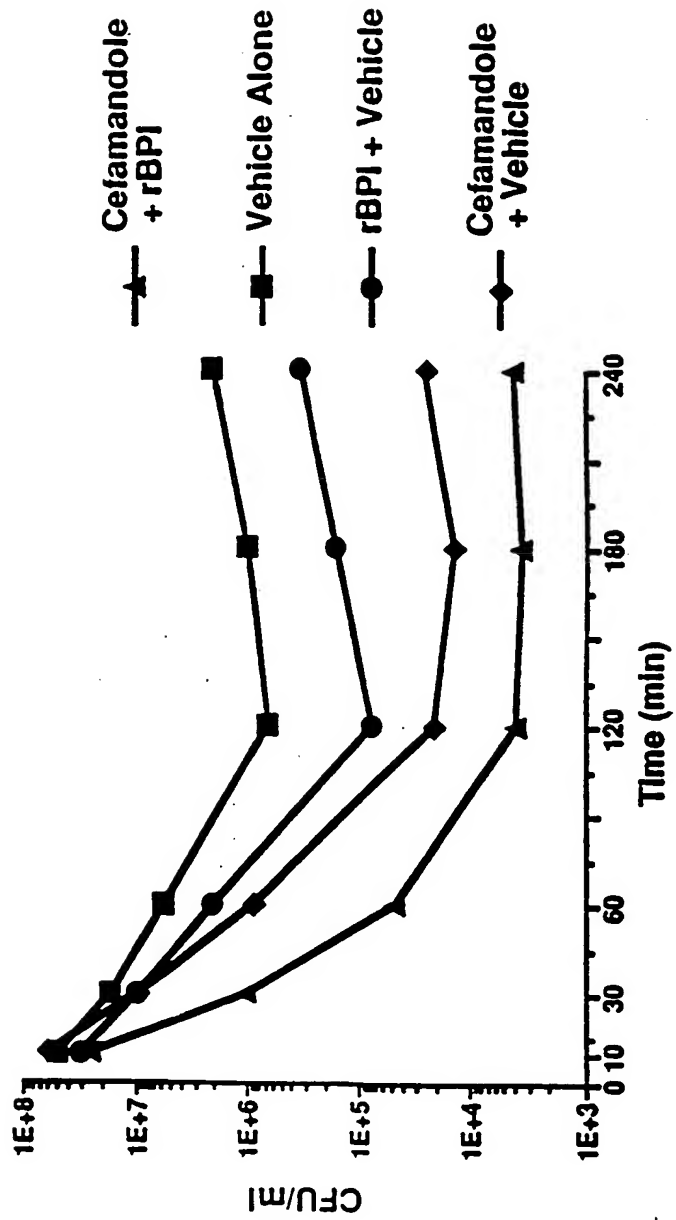


Fig. 7

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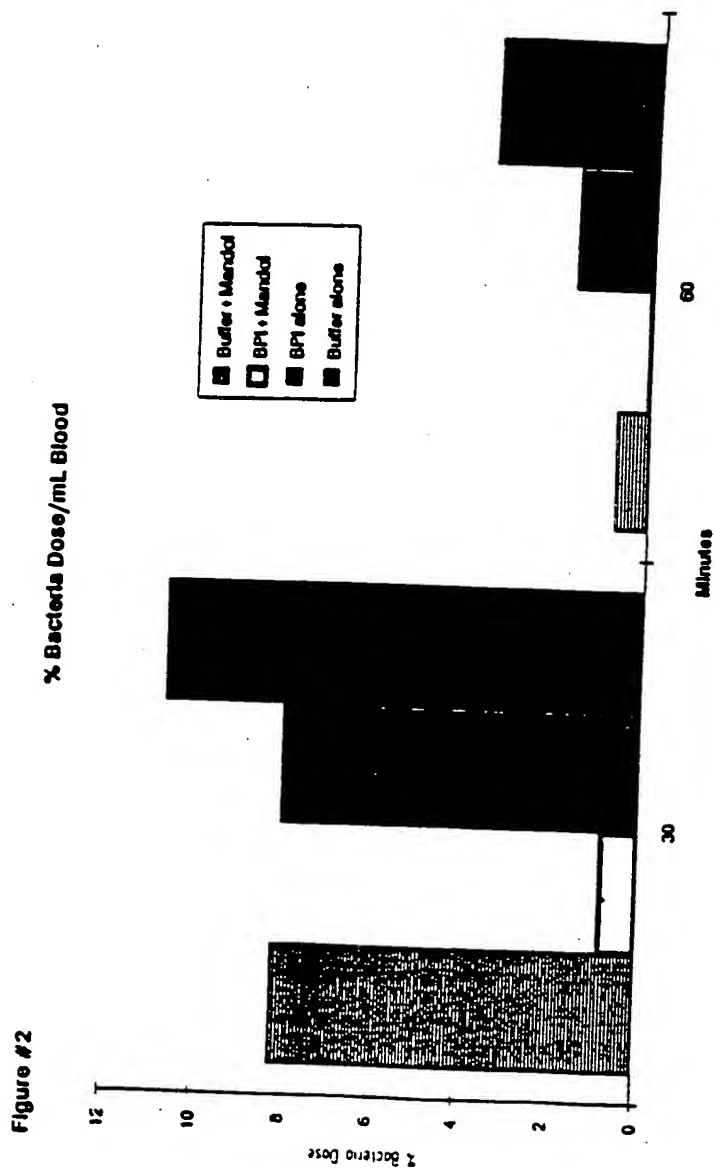


Fig. 8

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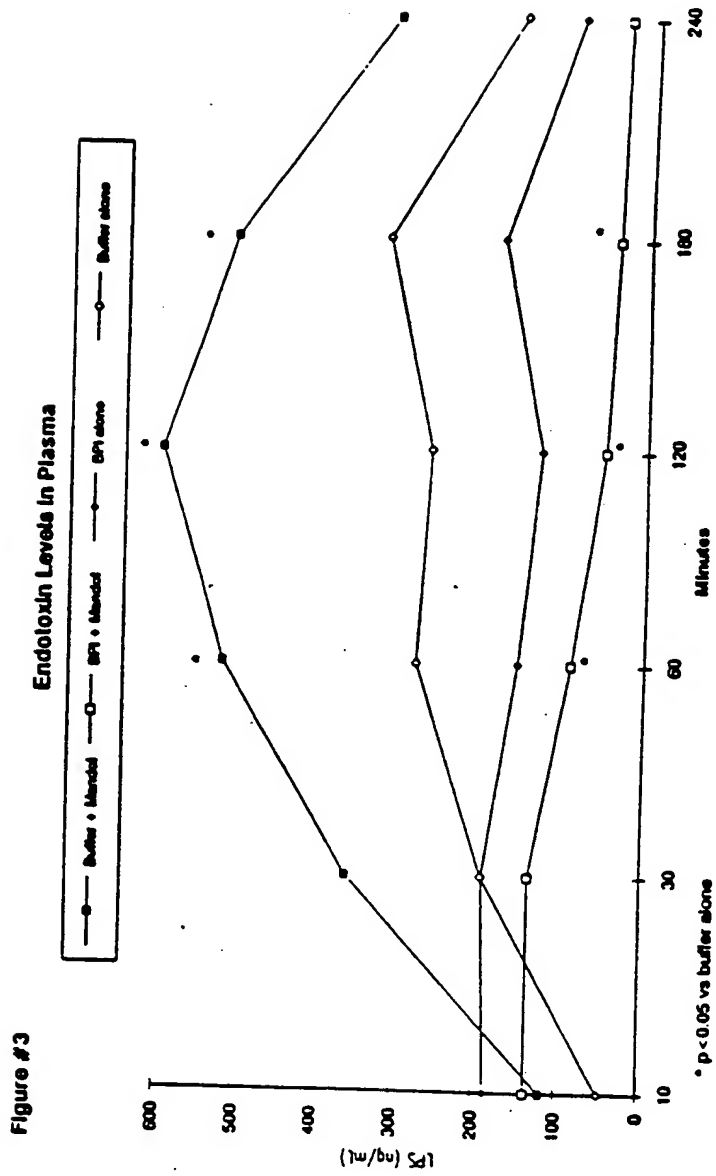


Figure #3

Fig. 9

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FIG. 10

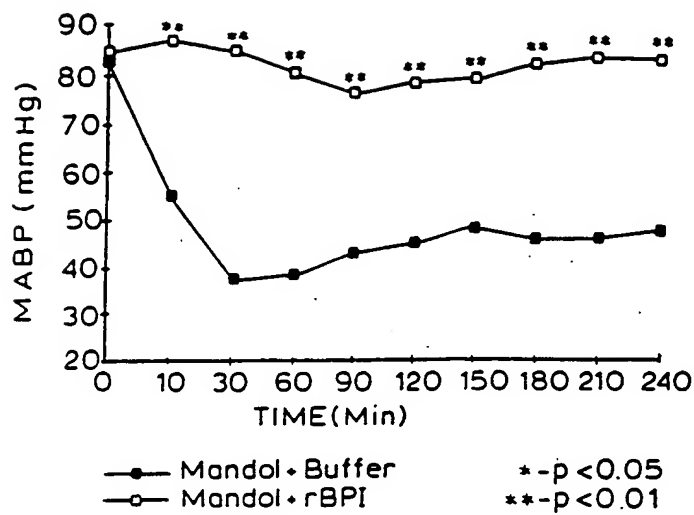
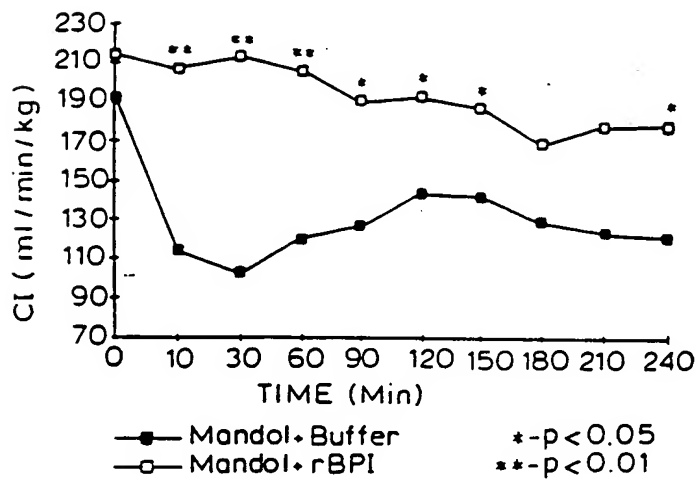


FIG. 11



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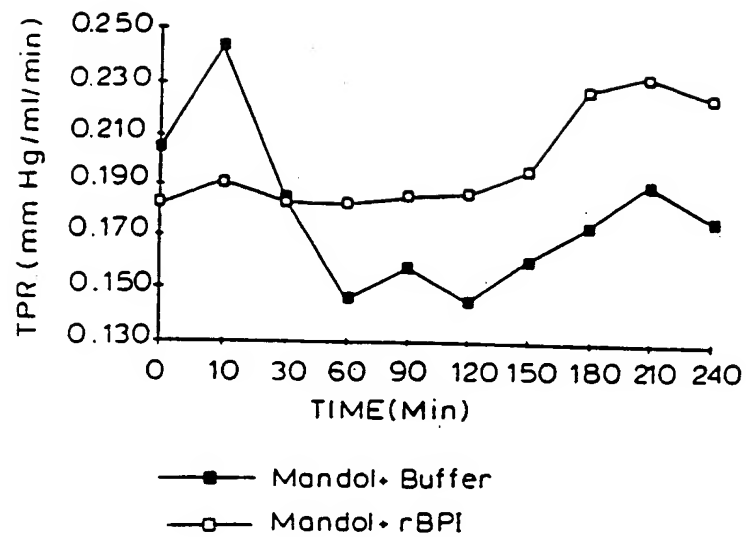


FIG. 12

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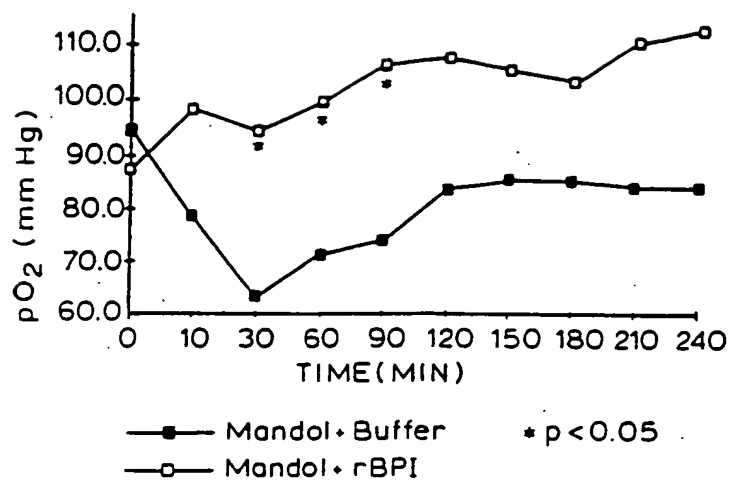


FIG. 13

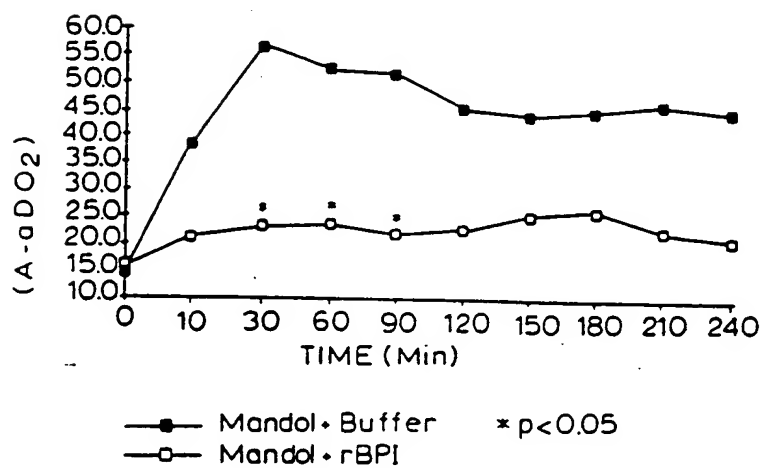


FIG. 14

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FIG. 15

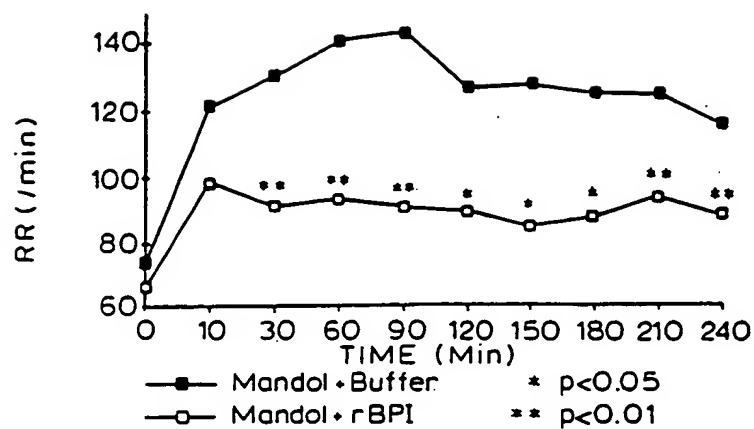
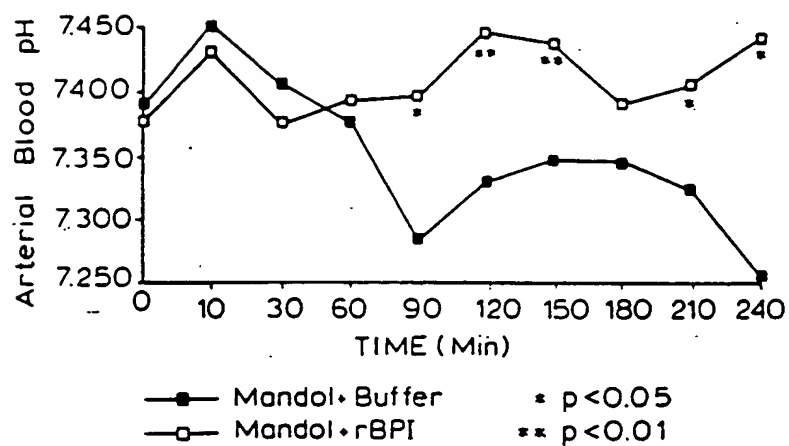


FIG. 16



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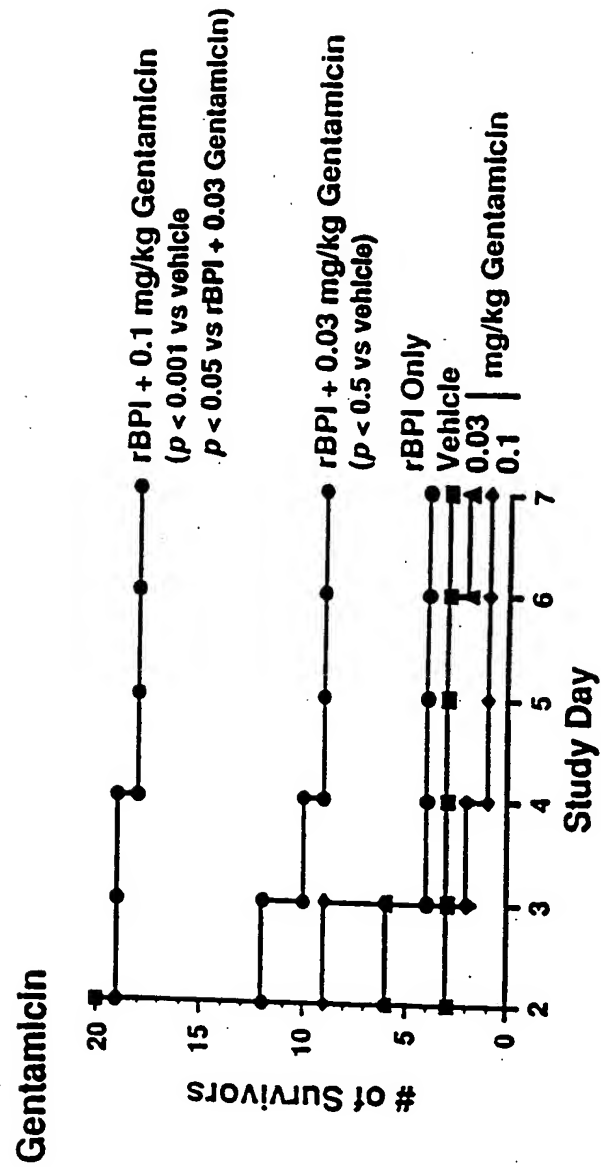


Fig. 17

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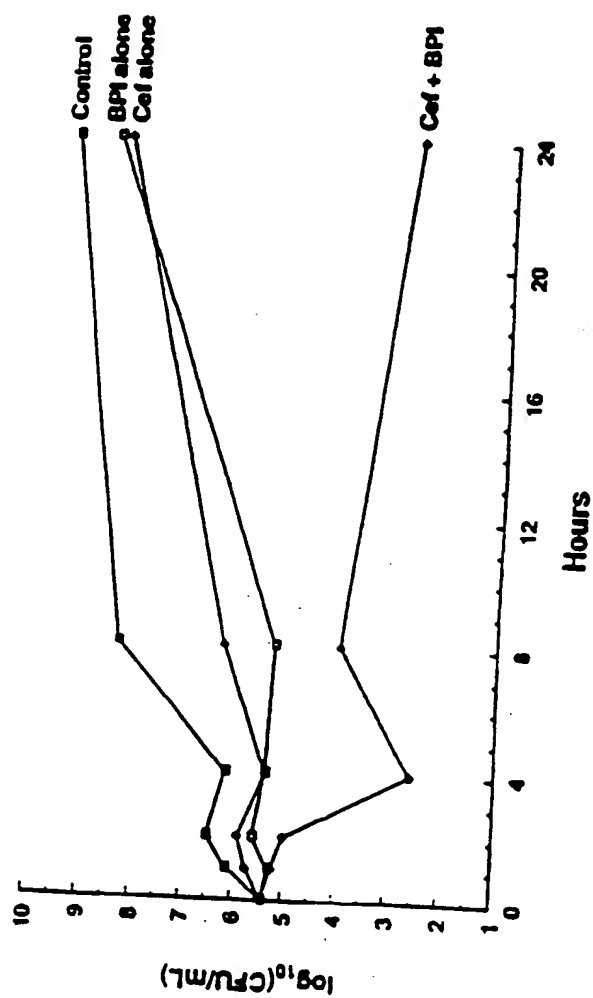


Fig. 18

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Fig. 19

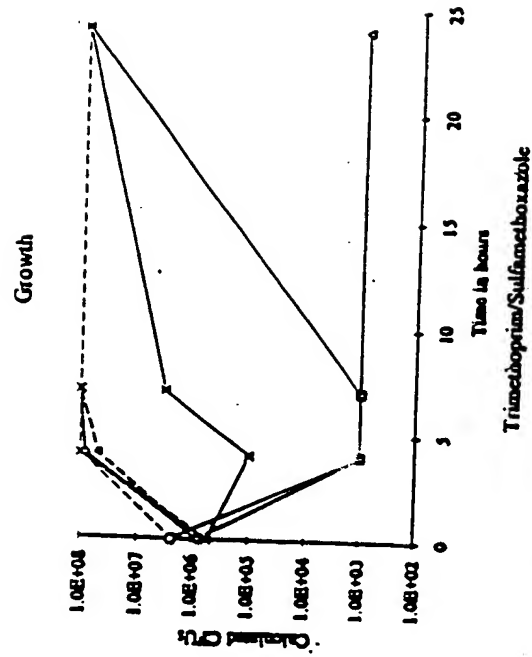
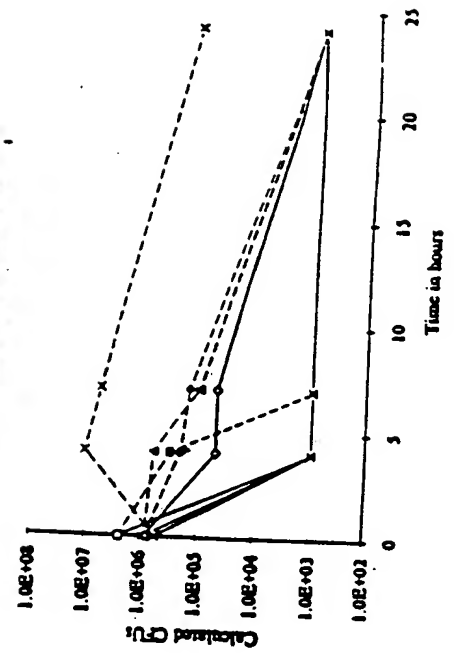


Fig. 20



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Fig. 21

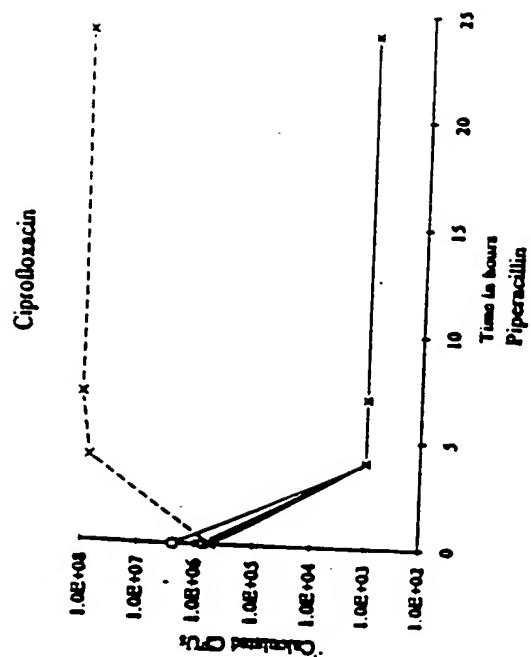
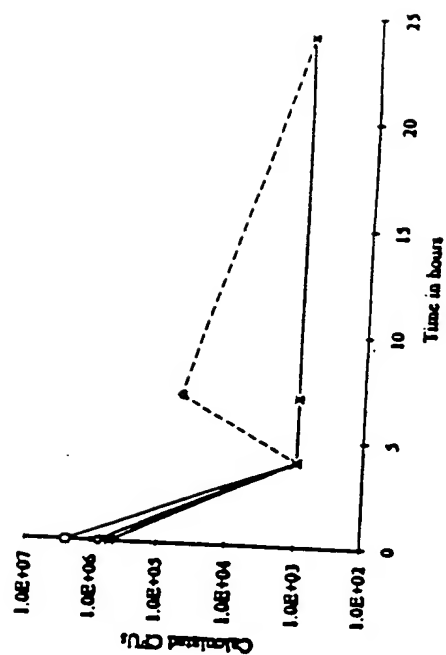


Fig. 22



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Fig. 23

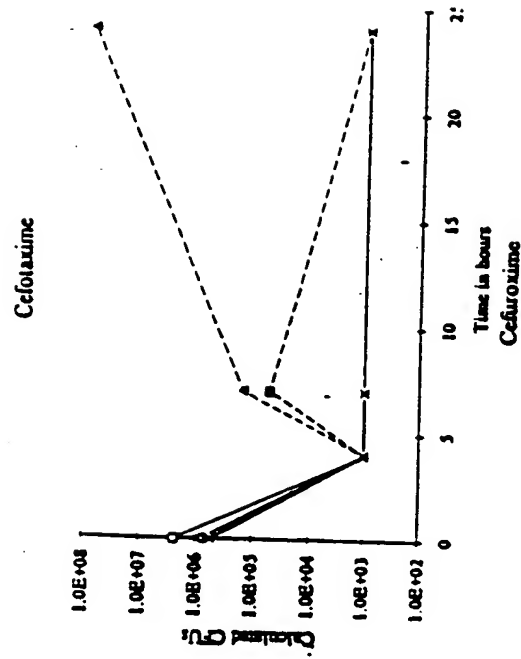
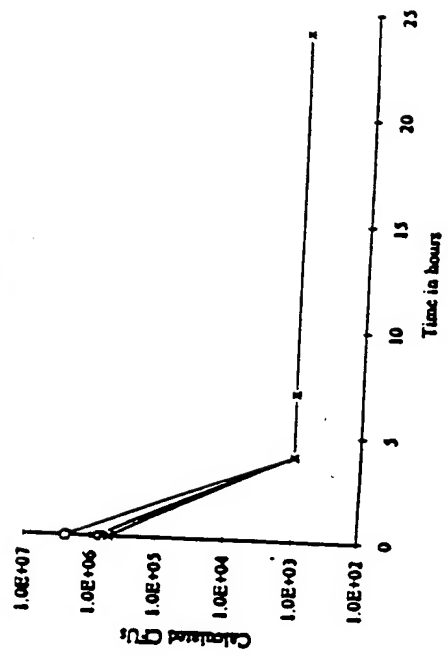
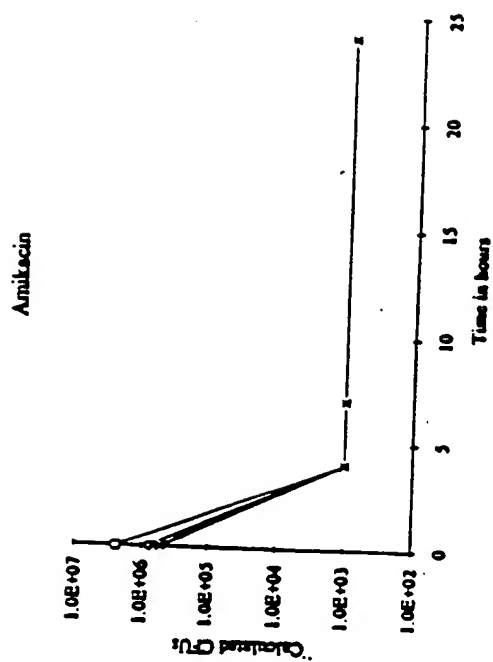


Fig. 24



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Fig. 25



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INTERNATIONAL SEARCH REPORT

International Application No
PCT/US 94/11225

A. CLASSIFICATION OF SUBJECT MATTER
IPC 6 A61K38/17

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
IPC 6 C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO,A,92 09621 (NEW YORK UNIVERSITY) 11 June 1992 see page 6, line 32 - page 7, line 2 see page 16, line 6 - page 17, line 19 see page 24, line 21 - page 27, line 28 see page 30, line 25 - page 31, line 9 ---	1-66
X	WO,A,89 01486 (NEW YORK UNIVERSITY) 23 February 1989 cited in the application see page 5, line 3 - line 7 see page 9, line 17 - page 12, line 8 see page 18, line 21 - page 21, line 22 see page 24, line 16 - line 35 -----	1-66

☐ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

* Special categories of cited documents:

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier document but published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

"A" document member of the same patent family

Date of the actual completion of the international search

7 February 1995

Date of mailing of the international search report

22.02.95

Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentaan 2
NL - 2230 HV Rijswijk
Tel. (+ 31-70) 340-2040, Tx. 31 651 epo nl,
Fax (+ 31-70) 340-3016

Authorized officer

Rempp, G

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US94/11225

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:
Remark: Although claims 33-37, 39-43, 45-49, 51-54, 56-60, 62, 65 and 66 are directed to a method of treatment of the human/animal body the search has been carried out and based on the alleged effects of the compound/composition
2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

Information on patent family members

Insert International Application No

PCT/US 94/11225

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
WO-A-9209621	11-06-92	AU-A- 9127591 EP-A- 0563222	25-06-92 06-10-93
WO-A-8901486	23-02-89	EP-A- 0375724 JP-T- 2504515 US-A- 5198541	04-07-90 20-12-90 30-03-93

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